Dysregulated Wnt Signalling in the Alzheimer’s Brain

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Abstract: The Wnt signalling system is essential for both the developing and adult central nervous system. It regulates numerous cellular functions ranging from neurogenesis to blood brain barrier biology. Dysregulated Wnt signalling can thus have significant consequences for normal brain function, which is becoming increasingly clear in Alzheimer’s disease (AD), an age-related neurodegenerative disorder that is the most prevalent form of dementia. AD exhibits a range of pathophysiological manifestations including aberrant amyloid precursor protein processing, tau pathology, synapse loss, neuroinflammation and blood brain barrier breakdown, which have been associated to a greater or lesser degree with abnormal Wnt signalling. Here we provide a comprehensive overview of the role of Wnt signalling in the CNS, and the research that implicates dysregulated Wnt signalling in the ageing brain and in AD pathogenesis. We also discuss the opportunities for therapeutic intervention in AD via modulation of the Wnt signalling pathway, and highlight some of the challenges and the gaps in our current understanding that need to be met to enable that goal.

Keywords: Wnt signalling; Alzheimer’s disease; neurodegeneration; APP processing; tau pathology; synapse degeneration; neuroinflammation; blood brain barrier

1. Introduction

The term dementia encompasses a group of devastating disorders with characteristic declines in cognition, function and behaviour, which eventually severely impair patients’ ability to perform instrumental and/or basic activities [1,2]. Beyond the patient, this profoundly impacts caregivers, families and society as a whole, which is facing ever increasing health care costs associated with dementia [3]. Alzheimer’s disease (AD) is by far the most common form of dementia accounting for 60–80% of cases [3], with an estimated 40 million diagnosed patients worldwide [4]. Current therapeutic options are restricted to symptomatic treatments such as cholinesterase inhibitors or the glutamate receptor antagonist memantine, which have a limited effect on memory and cognition [5]. As disease-modifying therapies, which target the disease process, are not yet available [6,7], AD dementia inevitably results in death within 5–12 years of symptom onset [8].

While the need for disease-modifying therapies that prevent disease onset or slow disease progression is clear, the lack of such treatments reflects the complicated, multifactorial pathobiology of AD. Patients often present with a range of disease manifestations, but what uniquely defines AD is the presence of protein aggregates in the form of extracellular deposits of β-amyloid (Aβ) as diffuse and neuritic plaques, as well as intracellular neurofibrillary tangles (NFTs) consisting of hyperphosphorylated tau [9] observed in post-mortem tissue. Yet, as of writing, therapeutic approaches targeting these two disease hallmarks have not proven to be effective in clinical trials [6,7] (although
approaches targeting tau are at a relatively early stage of evaluation in patients [10]). This is likely to be for a number of reasons, beyond the scope of this review. However, it does raise the possibility that an effective therapeutic strategy will require the targeting of several pathological manifestations via different biological pathways, and that this may change over the time course of the disease.

The Wnt signalling system plays a crucial role in many cellular processes such as cell differentiation, migration, and tissue homeostasis. In the central nervous system (CNS), Wnt signalling regulates developmental programmes and, as is increasingly recognised, it modulates a number of aspects of the mature brain such as synapse number and function, the integrity and function of the blood brain barrier (BBB), as well as the biology of microglia (the resident immune cells of the CNS). These key facets of the mature brain are significantly impacted in AD, raising the possibility that dysregulation of Wnt signalling may play an important role in a number of different aspects of this complex disease [11,12].

In this review, we aim to provide a systematic overview of the role of Wnt signalling in the CNS, together with what is known regarding dysregulation of this signalling system and how this contributes to certain pathological processes in AD. We also consider potential opportunities for new therapeutic approaches in AD through selective modulation of Wnt signalling, while recognising some key gaps in our knowledge that need to be addressed.

2. Wnt Signalling Pathways—An Overview

Wnt ligands are lipid-modified [13], secreted glycoproteins, which, upon binding to cell surface receptors, trigger intracellular signalling pathways that regulate various biological processes such as the cell cycle, cell migration and establishment of cell polarity [14,15]. The genomes of most mammals harbour 19 Wnt genes that can be grouped into 12 conserved subfamilies. These ligands are recognised by a heterodimeric receptor complex on the cell surface comprising Frizzled (Fz) as well as single-pass transmembrane co-receptors LRP5, LRP6, Ror1, Ror2, or Ryk proteins (Figure 1) [16–19]. The mammalian genome harbours 10 Fz genes, which encode 7-transmembrane (7TM) receptors that exhibit an N-terminal, extracellular, large cysteine-rich domain (CRD) used for Wnt binding [20–22]. However, ligand-receptor interactions are promiscuous as evidenced by the fact that there are multiple, non-mutually exclusive ligand receptor combinations [23,24].

Wnt ligand binding by the receptor complex induces conformational changes of the receptor complex and phosphorylation of target proteins as intracellular signalling pathways are initiated. These signalling pathways can broadly be classified into the canonical Wnt/β-catenin and non-canonical β-catenin-independent signalling cascades. The activation of specific pathways depends on, among various other factors, the exact identity of involved ligand and receptor isoforms, the expression of which is under tight spatiotemporal control [23].

The canonical Wnt/β-catenin signalling cascade is initiated upon binding of Wnt to the Fz-LRP receptor complex (Figure 1a,b) [25]. This subsequently leads to the inactivation of a multiprotein complex consisting of CK1α, GSK-3β, Axin, and APC, which usually phosphorylates β-catenin, thus marking it for proteasomal degradation [26]. Accumulating β-catenin can now translocate to the nucleus where it associates with transcription factors from the TCF/Lef family [27,28] to activate transcription of Wnt/β-catenin target genes with known roles in proliferation, fate specification and differentiation in development as well as adult tissue homeostasis [29].

Initiation of the non-canonical Wnt/PCP (planar cell polarity) cascade requires Wnt binding to a Fz-Ror/Ryk receptor complex (Figure 1c) [30–32]. Intracellularly, this causes the activation of small GTPases RhoA, Rac, and Cdc42 [33,34], which subsequently activate the downstream kinases JNK and ROCK [35,36], which in turn regulate actin and microtubule cytoskeletons. As such, the Wnt/PCP cascade plays a vital role in the control of cell/tissue polarity and cell migration [37].
Wnt-activated inhibitory factor 1 (Waif1) There are seven secreted antagonist families (the Dickkopf proteins (Dkks), secreted Frizzled-related proteins (sFRPs), Wnt inhibitory factor 1 (WIF-1), Wise/SOST, Cerberus, insulin-like growth factor binding protein 4 (IGFBP-4), Notum and four transmembrane Wnt antagonist families (Shisa, Wnt-activated inhibitory factor 1 (Waif1/5T4), adenomatosis polyposis coli down-regulated 1 (APCDD1),

Figure 1. Wnt signalling pathways. Note that emphasis is placed on pathways and components that are of greater relevance in the context of AD. (a) Canonical Wnt signalling pathway, off: In absence of Wnt ligand binding to Frizzled (Fz)-LRP5/6 heterodimeric cell surface receptors, the Axin/APC/CK1α/GSK-3β destruction complex phosphorylates the transcription factor β-catenin, marking it for proteasomal degradation. Endogenous canonical Wnt antagonists (including sFRP and Dkk1) further favour the ‘off’ state. (b) Canonical Wnt signalling pathway, on: Wnt ligand binding to Fz-LRP5/6 recruits the scaffolding protein DVL, which in turn inhibits GSK-3β activity of the destruction complex. β-catenin subsequently accumulates in cytoplasm, allowing it to translocate to the nucleus where it transactivates the TCF/LEF-mediated expression of canonical Wnt target genes. (c) Wnt/planar cell polarity (PCP) pathway: Wnt ligand binding to Fz-ROR/Ryk co-receptors recruits DVL, which in turn activates the small GTPases RHOA, RAC, and CDC42. RHOA and RAC conjoinly activate JNK, which stimulates Jun transcription factor dependent gene expression via phosphorylation. Furthermore, RHOA, via ROCK, and CDC42 stimulate actin cytoskeleton remodelling.

The Wnt/Ca²⁺ cascade requires binding of Wnt ligand to Fz receptor, which intracellularly triggers G-protein coupled signalling [38]. This in turn activates phospholipase C (PLC) [39], which stimulates the release of Ca²⁺ from intracellular stores via the signalling molecule inositol triphosphate (IP3) [40]. The mobilised Ca²⁺ then stimulates the Ca²⁺-sensitive protein kinases protein kinase C (PKC) [41] and Ca²⁺/Calmodulin-dependent protein kinase II (CaMKII) [42], as well as the Ca²⁺-sensitive transcription factor NF-AT. Through these effector proteins the Wnt/Ca²⁺ cascade regulates many processes, ranging from developmental cell fate determination, cell/tissue migration, cell differentiation, and inflammatory response mediation [43].

Wnt signalling can furthermore be modulated by a number of endogenous agonists and antagonists [44], which are important for the fine-tuning of Wnt signalling-regulated processes. There are seven secreted antagonist families (the Dickkopf proteins (Dkks), secreted Frizzled-related proteins (sFRPs), Wnt inhibitory factor 1 (WIF-1), Wise/SOST, Cerberus, insulin-like growth factor binding protein 4 (IGFBP-4), Notum and four transmembrane Wnt antagonist families (Shisa, Wnt-activated inhibitory factor 1 (Waif1/5T4), adenomatosis polyposis coli down-regulated 1 (APCDD1),
and Tiki1). They exert their function either by sequestering/inactivating secreted Wnt (e.g., WIF1, Cerberus, sFRP, Notum) or by blocking/sequestering elements of receptor complexes (e.g., Dkk1, Wise/SOST, IGFBP-4). For example, Dkk1 sequesters LRP6, thus preventing its heterodimerisation with Fz8 to block canonical Wnt/β-catenin signalling [45]. When the Dkk1 co-receptor Kremen 2 (Krm2) is present, this additionally leads to the endocytosis of LRP5/6-Krm2-Dkk1 complexes [46]. The related Dkk2, however, can act both as an activator and as an inhibitor of the Wnt/β-catenin cascade [47], in the presence Krm2, Dkk2 functions as an LRP6 antagonist, while in its absence it functions as an activator.

Conversely, there are two families of secreted proteins that act purely as Wnt agonists: R-spondins (Rspo) and Norrin. R-spondins stimulate canonical Wnt/β-catenin signalling by promoting the internalisation of the transmembrane E3 ubiquitin ligase ZNRF3 that usually marks Fz and LRP6 for degradation [48]; this is mediated by the Rspo receptors Lgr4, Lgr5, and Lgr6 [49]. R-spondins can also stimulate Wnt/PCP signalling via a mechanism that requires Wnt5a-Fz7 signalling, Rspo3 binding to the four transmembrane proteoglycan syndecan 4, and syndecan 4-dependent endocytosis of the entire Wnt5a-Fz7-Rspo3-syndecan 4 complex [50]; in contrast to the action of Dkk1/Krm2 on canonical Wnt/β-catenin signalling through Fz8/LRP6, this internalisation was shown to be crucial for Wnt/PCP signal transduction. Finally, Norrin, although structurally unrelated to Wnts, binds Fz4/LRP5 to activate Wnt/β-catenin signalling [51].

The Wnt cascades thus constitute a highly complex signalling network with a range of different functions and roles in development, mature homeostasis, ageing, as well as disease. The remainder of this review will provide more detailed insight into Wnt signalling in the brain with a focus on the deregulation of Wnt signalling in AD.

3. Wnt Signalling in the Brain

Wnt signalling plays an important role in various aspects of the brain, ranging from brain development to normal brain function. Indeed, the gene expression of various Wnt and Fz receptor isoforms is subject to tight spatio-temporal control, as has been reviewed in [52]. Unsurprisingly, altered Wnt signalling strength can have detrimental effects on the brain, as is for instance observed during ageing where reduced Wnt signalling is evident. In this section we summarise the importance of Wnt cascades within the healthy brain lifespan.

3.1. Wnt Signalling during Brain Development

Wnt signalling plays an important role during various aspects of brain development. As secreted morphogens, Wnts as well as modulators of Wnt signalling, regulate the establishment of distinct tissue domains within the CNS along Wnt concentration gradients leading to the specification of the brain as a whole and substructures within the brain in a process that is more generally termed tissue patterning. Important Wnt signalling components in this process include Wnt-1, Wnt3a, Wnt8c, Dkk1, sFRPs, and Notum [53–59]. Fittingly in the context of AD, Wnt signalling is also required for the development of brain structures including the hippocampus among others, which is integral for memory [60]. There, Wnt3a regulates both neuronal progenitor cell (NPC) expansion and neuronal differentiation via canonical signalling [61–65]. Wnt3a is further accompanied by Wnt5a and Wnt7b, which respectively regulate axonal and dendritic differentiation (axon specification, axon growth, and dendritic tree formation) via non-canonical Wnt/PCP signalling [66–70]. One further neuronal differentiation aspect also regulated by Wnt signalling is synaptogenesis, as has been reviewed in [52]. Canonical Wnt signalling (via Wnt7a, Wnt7b, and Wnt3a) and non-canonical Wnt signalling (via Wnt5a) respectively increased and decreased pre-synaptic assembly in rat primary hippocampal neurons [71–75], indicating that canonical and non-canonical Wnt pathways need to be well balanced to ensure adequate synapse numbers (this concept is relevant in the context of AD, as will be described in Section 4.3). Conversely, post-synaptic assembly is driven by Wnt5a-mediated non-canonical Wnt/PCP as well as Wnt7a-mediated Wnt/Ca2+ pathways, at least in rat primary hippocampal neurons [76–79]. Taken together, Wnt signalling is highly important in the brain at many stages of development.
3.2. Wnt Signalling in the Mature Brain

Beyond the developmental period, Wnt signalling remains integral to brain function in adulthood, especially at the level of the synapse. Wnts and Wnt regulatory proteins are released locally at the synapse. This ensures homeostatic control over synaptic connectivity even during basal synaptic transmission. A study on mature cultured rat hippocampal neurons and adult mouse brain slices indicates the presence of an endogenous, non-canonical Wnt5a signalling tone that regulates excitatory synapse activity [76]; blocking endogenous Wnt signalling by exogenous sFRP antagonist treatment reduced the amplitudes of evoked field excitatory post-synaptic potentials (fEPSPs) and miniature excitatory post-synaptic currents (mEPSCs) according to electrophysiology recordings. In mature mouse brain slices containing the entorhinal cortex-hippocampal circuit, antibodies against canonical ligands (Wnt3a and Wnt7a) and non-canonical ligand (Wnt5a), as well as application of sFRP-2 reduced circuit activity; this led the authors to conclude that constitutive Wnt release contributes to network maintenance, presumably by affecting synaptic mechanisms [80]. Among the two glutamate receptors present at the post-synaptic terminus of excitatory synapses, α-amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid receptor (AMPAR) and N-methyl-D-aspartate receptor (NMDAR), Wnt5a scavenging with sFRP treatment specifically reduced basal NMDAR, but not AMPAR, currents (Wnt5a usually increases NMDAR currents in a Ca\textsuperscript{2+}, PKC, and JNK dependent manner) [81]. Another sFRP treatment study revealed that Wnt signalling regulates basal synaptic transmission by maintaining pre-synaptic termini in the CA3 stratum lucidum of the hippocampus, which receives axonal inputs from the dentate gyrus (DG) area of the mouse hippocampus [82]. Taken together, these studies suggest that a basal endogenous Wnt signalling tone is required to ensure structural and functional synaptic connectivity.

Beyond basal synaptic transmission, Wnt signalling pathways also refine synaptic contacts (strengthening or weakening) in an environment and activity-dependent manner—a mechanism known as synaptic plasticity [83]. In cultured rodent hippocampal neurons and slices, Wnt7a increased the density and maturity of dendritic spines, and further enhanced synaptic strength by AMPAR recruitment to synaptic spines in a CaMKII-dependent manner [79,84], while simultaneously enhancing neurotransmitter release at CA3-CA1 synapses [72,85]. Protocols that elicit long term potentiation (LTP; the sustained strengthening of a synapse thought to be the cellular model of learning and memory), stimulate Wnt3a synaptic secretion upon post-synaptic NMDAR activation in mice [86]. Newly released Wnt3a subsequently binds synaptic Fz receptors to canonically initiate the expression of LTP target genes. Electrophysiologically, canonical Wnt3a enhanced the LTP response by increasing fEPSPs in adult mouse hippocampal slices [86]. At least in rat and mouse hippocampal slices, this appears to be complemented by non-canonical Wnt5a signalling, which, via NMDAR current potentiation, facilitated the induction of LTP [81,87]. The above findings were also supported by an in vivo study, which demonstrated that environmental enrichment increased Wnt7a/b levels in the mouse hippocampus; this in turn induced synaptic remodelling in the CA3 of the mouse hippocampus, suggesting that Wnt signalling can bring about structural changes even in the adult brain [82]. Furthermore, spatial information storage was also correlated with increased Wnt7 and Wnt5a expression in the mouse DG 7 days and 30 days after water maze training [88]. Indeed administration of activators of canonical (WASP-1) and non-canonical signalling (FOXY-5) improved episodic memory in a water maze spatial task [87]. A subsequent study provided further proof that canonical Wnt signalling is necessary for hippocampal memory consolidation in mice, as canonical Wnt signalling was enhanced following object recognition training, which could be blocked by hippocampal injection of Dkk1 [89].

Neurogenesis, which in the adult brain is restricted to a select few brain regions including the subgranular zone (SGZ) of the DG, is also modulated by Wnt signalling. Neurogenesis has been reported to influence memory [90], though how exactly adult neurogenesis contributes to memory and whether it even exists in humans is still a matter of ongoing debate [91]. Nevertheless, research in rats has shown that Wnt3-mediated canonical signalling stimulates adult neurogenesis in the DG in vitro and in vivo, and that loss and gain of function approaches to block or stimulate Wnt signalling
in the DG respectively attenuated and increased adult neurogenesis [92,93]. Furthermore, blocking Wnt signalling in the DG of adult rats impaired spatial and object recognition memory [94]. Wnt3a canonical signalling was shown to transcriptionally induce expression of NeuroD1 in mice, which has a role in the neuronal differentiation of DG-resident granule cell neurons [95,96]. In addition to Wnt3, canonical Wnt7a signalling may also play a role in adult neurogenesis in the mouse DG, where a decreased NPC population, reduced neurogenesis, and impaired granule cell neuronal differentiation were evident in Wnt7a knockout mice compared to control; these results indicated that Wnt7a regulates both NPC proliferation (via cyclin D1) and neuronal differentiation (via Neurog2) [97]. The endogenous Wnt inhibitor sFRP3 also regulates both NPC proliferation and neuronal differentiation, where adult sFrp3 knockout mice exhibit both increased NPC proliferation, neurogenesis, and accelerated dendritic growth and spine formation of new-born neurons [93]. A similar negative regulatory function was recently described for endogenous Wnt antagonist Notum, which limits adult olfactory bulb neurogenesis from the subventricular zone in mice [98]. Taken together, Wnt signalling is crucial for the regulation of neuronal circuits, both at a baseline level as well as during activity-related plastic changes that likely contribute to memory and learning.

3.3. Wnt Signalling in the Ageing Brain

Ageing is a multifactorial process of progressive detrimental changes occurring at the molecular, cellular, tissue, and organ level and is associated with functional decline [99]. This also applies to the ageing brain, where impairments in cognitive function and memory are evident. A body of work suggests that this might be due reductions in both synapse function and adult neurogenesis, particularly in the hippocampus [100–103]. Interestingly, dysregulated Wnt signalling is also observed in older age. Indeed, several genes involved in Wnt signalling are downregulated in the brains of advanced age rodents, pointing towards reduced Wnt signalling tone. In 36 months old mice, the Wnt ligands Wnt2, Wnt4, and Wnt8a, as well as the canonical transcription factors Lef1, Tcf3, and the transcriptional output Ccnd2 were significantly decreased compared with 5 month controls, as determined by quantitative PCR [104]. Meanwhile, astrocytic Wnt3a expression and secretion in the DG progressively declined, while Dkk1 expression increased with age [105,106]. In a comparison between 12 or 24 month old rats with 4 month old controls, reduced total as well as nuclear β-catenin protein were detected in the hippocampus alongside increased phosphorylated (active) GSK-3β, indicative of diminished canonical Wnt signalling in aged animals [107]. The aged human brain also exhibits gene expression patterns that are in line with reduced Wnt signalling, specifically a downregulation of the Wnt ligands WNT2B, WNT6, WNT7a, and Fz receptors FZD2 and FZD3, alongside an upregulation of GSK3β and SFRP1 in the prefrontal cortex [108]. Further evidence implies that menopause-related estrogen deprivation increases DKK1 expression in the mammalian hippocampus [109,110].

Given the well documented role that Wnt signalling plays in the healthy, mature brain many studies now suggest that the Wnt signalling-dampened aged brain is prone to reductions in both synapse function and adult neurogenesis, thus contributing to cognitive decline. Indeed, conditional Dkk1 knock out in NPCs restored adult neurogenesis in old age mice, and increased dendritic complexity as well as neuronal activity in the DG [106]. This concomitantly restored working memory and memory consolidation in behavioural tests. Conditional Dkk1 overexpression in mature 3-6 months old mice (thus modelling Dkk1 upregulation at more advanced ages) had profound consequences on synapses, resulting in excitatory synapse loss, reduced LTP, and learning/memory deficits [111]. Crucially, this was fully reversible upon cessation of Dkk1 overexpression. In another study, age-related decline in astrocytic Wnt3a expression and secretion reduced adult neurogenesis in mice; however, this could be reversed by overexpressing Wnt3a in astrocytes or by stimulating endogenous astrocytic Wnt3a production with exercise [105]. Finally, stimulation of canonical and non-canonical Wnt signalling with small molecule “activators” FOXY-5 and WASP-1, improved excitatory synapse transmission, LTP and episodic memory in mature mice [87].
The underlying tenor is therefore that Wnt signalling strength declines with increasing age and that this can lead to cognitive decline. Equally, restoring deficient Wnt signalling may be able to reverse age-related cognitive deficits, although the strategy of how this can be effectively achieved in humans remain to be elucidated (see Section 4.6). As discussed in Section 5, any future ambitions to enhance Wnt signalling using small molecules or gene therapy approaches requires careful consideration with respect to the exact Wnt pathway target, cellular specificity, and dose so as to avoid unwanted side effects including tumour formation.

4. Wnt Signalling in AD

Given the connection between age-related cognitive decline and decreased Wnt signalling strength (discussed in the previous section), it is perhaps unsurprising that Wnt signalling pathways are also suppressed in AD given the strong connection between AD and its biggest risk factor–age [112,113]. Indeed, the majority of AD cases are detected at an advanced age (sporadic late onset AD or LOAD) supporting the fact the AD is predominantly an age-related disorder of the brain. However, mutations in certain genes result in early onset heritable/familial AD or FAD; see Section 4.1). While FAD is associated with mutations in amyloid precursor protein (APP) or APP processing genes (PSEN1, PSEN2, encoding components of the γ-secretase complex), numerous additional risk genes (including APOE, TREM2, and UN5C5) as well as environmental factors confer susceptibility to LOAD. Genetic studies have been invaluable in shedding light on the multiple pathological mechanisms observed in AD, as will be discussed below. In several cases where gene variants have been linked to AD, these genes also show connections to Wnt signalling pathways.

Among the first pieces of evidence indicating that abnormal Wnt signalling could be involved in AD came from a study that discovered a polymorphism in the GSK3β promoter, which increased its activity and conferred increased susceptibility to LOAD [114] (given the known biological role of GSK-3β, increased activity reduces canonical Wnt signalling by promoting β-catenin degradation). Further supporting the link between reduced canonical signalling and AD, a single nucleotide polymorphism (Ile-1062 → Val) and a novel splice variant within the canonical Fz co-receptor encoding gene LRP6 were also associated with LOAD; functionally, in HEK293T cells, both reduced canonical Wnt signalling [115,116]. In addition, LRP6 mRNA and protein levels were significantly decreased in human AD brains compared to controls [117] (for an extensive review on the connection between LRP6 and AD the reader is referred to [118]). Another Wnt signalling suppressing component in AD is the induced neuronal expression of the endogenous Wnt signalling antagonist Dkk1 in the brains of post-mortem AD patients as well as in AD mouse models [119,120]. In fact, it was demonstrated that AD-associated Aβ fibrils induce Dkk1 expression in acute mouse hippocampal slices within several hours [121]. Importantly, DKK1 is a ligand for LRP6 and exerts it action by removing LRP6 from functional canonical Fz-LRP6 receptor heterodimers [45]. Interestingly, the Dkk1 homologue Dkk2 was also significantly upregulated in various AD mouse models, specifically within the myeloid cell lineage that includes microglia and other immune cells [122]. The protein clusterin also participates in the Aβ-DKK1 pathway, as discussed below in Section 4.3. From a purely genetic standpoint, the encoding gene CLU had previously been identified as a major LOAD risk gene in genome wide association studies (GWAS), where various AD-linked single cell nucleotide polymorphisms were found to be associated with AD [123,124].

ApoE is a regulator of lipid homeostasis predominantly produced by astrocytes within the CNS. By binding to ApoE receptors on neurons, it transports cholesterol to neurons, which lack cholesterol producing capabilities. [125,126]. However, among the three existing polymorphic alleles ε2, ε3 and ε4, the APOE ε4 allele is found in ~40% of AD cases despite only being represented in 13.7% of the general population, thus classifying it as a major AD risk gene [127]. The nature of the role of ApoE in AD is multifaceted and still not fully understood, but highlighted by increased Aβ binding and deposition in APOE ε4 carriers [128]. In addition, collective evidence suggests that ApoE also interacts with Wnt signalling pathways in an AD-relevant manner: ApoE4-mediated Aβ pathology in the APP/PS1
AD mouse model required the expression of its neuronal receptor LRP1 [129], which had previously been found to suppress Wnt3a-driven canonical Wnt signalling in HEK cells by interacting with Fz1 [130]. Furthermore, ApoE4 treatment in PC12 cells, more so than ApoE2 and ApoE3, suppressed Wnt7a-stimulated canonical signalling [131].

Genetic variants of triggering receptor expressed on myeloid cells-2 (TREM2) have also been linked to increased susceptibility to LOAD [132,133]. Although relatively rare overall, carriers of the most common and best studied TREM2 variant (R47H) had a 2–4 fold increased risk for LOAD, which is in the same range as the APOE ε4 allele (minor allele frequency is population dependent but reaches up to 0.63% in the Icelandic population [132]). Within the CNS, TREM2 is predominantly expressed by microglia where it was shown to modulate canonical Wnt signalling to support microglial survival and microgliosis, both of which are markedly impaired in Trem2−/− mice [134]. These and other findings highlight the importance of the CNS immune system in AD, the proper function of which requires canonical Wnt signalling (this will be expanded upon in Section 4.4).

In addition to genetic analyses of AD pathogenic mechanisms, dynamic gene expression changes associated with AD and the resulting protein level changes are equally important for enabling our understanding of this disease. In a recent mass spectrometry based proteomics study, protein level changes were assessed in the brains and cerebrospinal fluid of AD patients and compared with control, prodromal, and mild cognitive impairment (MCI) cases [135]. Indeed, Wnt signalling related proteins were among those proteins exhibiting increased levels. These proteins included the Wnt ligands WNT5A, WNT5B as well as the endogenous antagonists SFRP1 and FRZB (SFRP3). While elevated SFRP1 and FRZB levels would be consistent with a reduced Wnt signalling tone, increased presence of WNT5A and WNT5B may reflect a pathological shift from canonical to non-canonical signalling that reduces synapse stability due to increased actin cytoskeleton dynamics [136] (see Section 4.3). In further support of a reduced Wnt signalling tone in AD brains, the phosphoproteome (phosphorylation status of proteins within the proteome) revealed significant phosphorylation increases of GSK-3β target proteins, which is indicative of increased GSK-3β activity and reduced canonical Wnt signalling strength [135]. These findings are corroborated by a further proteomics study, which reported that canonical Wnt signalling was dysregulated in specific human AD brain regions versus control [137]. The hippocampus for instance exhibited 17 differentially expressed Wnt signalling related proteins, including GSK-3β, GSK-3α, and AKT3, which were all downregulated, as well as the DKK homologue DKK3, which was upregulated. While the role of DKK3 as an agonist or antagonist of Wnt signalling is context dependent [138,139], the downregulation of GSK-3β, GSK-3α, and AKT3 would be expected to result in increased β-catenin stability and hence canonical Wnt signalling. This would be incompatible with the widespread notion that a reduction of canonical Wnt signalling contributes to AD pathology.

These findings notwithstanding, Wnt signalling dysregulation has been found to be a prominent feature in AD and further studies are needed to fully understand all the aspects involved. The following sections will provide a more detailed description of the role of Wnt signalling in the various pathobiological disease manifestations of AD (schematically summarised in Figure 2).
Figure 2. Wnt signalling in AD. Dysregulated Wnt signalling may contribute to various pathological manifestation of AD, including Aβ pathology, Tau pathology, neuronal pathology, synapse pathology, microglial pathology/neuroinflammation, as well as BBB pathology. Known or hypothesised (indicated by '?') changes in Wnt signalling associated with each manifestation are indicated (based on published literature). ↑ and ↓ indicate increased and decreased Wnt signalling respectively. Created with Biorender.com.

4.1. Wnt Signalling, APP Processing, and AD

One of the defining hallmarks of AD is the extracellular aggregation of Aβ and the formation of plaques. Its pathologic accumulation in the brains of AD patients is the result of aberrant proteolytic processing of the membrane–embedded Amyloid Precursor Protein (APP; APP family members usually play an important role during brain development and maintenance) (reviewed by [140]). Whereas the non-amyloidogenic processing pathway requires successive cleavage by α- and γ-secretase, the amyloidogenic pathway (the pathway resulting in the cleaved 39–42 amino acid long Aβ peptide (Aβ) relevant to AD) requires cleavage by β- and γ-secretases [141]. It is the pathological accumulation of Aβ caused by increased amyloidogenic processing and/or impaired Aβ clearance that contributes to
AD-related formation of Aβ plaques. While the 40 and 42 amino acid long peptides Aβ40 and Aβ42 are most prevalent in AD, it is Aβ42 that is particularly neurotoxic [141]. In keeping, mutations in genes involved in APP processing often lead to FAD. In fact, mutations in PSEN1 (presenilin 1 or PS1), encoding the catalytic subunit of γ-secretase, are the most common cause of FAD, followed by mutations in PSEN2 and APP [142–145].

The link between APP processing and impaired Wnt signalling can be exemplified by the fact that APP locus duplication causes FAD, involving a mechanism whereby APP binds β-catenin—as demonstrated in vitro, in vivo and in AD patients—preventing its nuclear translocation and thus canonical Wnt signalling [144,146]. β-catenin was also shown to interact with PS1; however, while this interaction usually stabilised β-catenin in cell lines, FAD-mutant PSEN1 exhibited reduced β-catenin stability and protein levels [147] and impaired trafficking to the nucleus [148] in HEK cells. This suggests that FAD-associated PSEN1 mutations negatively regulate canonical Wnt signalling. However, conflicting findings have been reported where human PSEN1 mutations increased steady state levels of β-catenin in neuronal (PC-12) and fibroblast cell lines (EcR293) [149,150]. While this is difficult to reconcile, it was suggested that these different outcomes may have been observed due to the use of different experimental models, different time points, the transient nature of the Wnt pathway involving negative feedback loops, measuring cytoplasmic rather than nuclear β-catenin, or potential cross-talk with the Notch pathway (the Notch receptor is a γ-secretase substrate and can context-dependently antagonise Wnt signalling by modulating β-catenin stability) (reviewed by [151]). Nonetheless, it remains highly likely that FAD-PSEN1 mutations, and by extension Aβ, are linked to dysregulated canonical Wnt signalling via dysregulation of β-catenin. Of note, reduced β-catenin levels were not observed in sporadic AD patients [147].

Thus, common FAD-associated variants of genes involved in APP processing can negatively impact Wnt signalling. However, the inverse is also true. Wnt signalling influences APP processing where inhibition of Wnt signalling was shown to induce amyloidogenic APP processing and Aβ aggregation while Wnt3a stimulation decreased the expression and activity of the amyloidogenic β-secretase BACE1 and amyloidogenesis in vivo and in vitro [152–154]. In further support, the canonical Wnt co-receptor LRP6 was found to be downregulated in human AD brains, which was correlated with a reduction of β-catenin and an increase of Aβ42 [117]. The same study then confirmed a causative relationship by showing that Lrp6 loss of function increased Aβ40 and Aβ42 production in neuronal cell lines, in the APP/PS1 AD mouse model, and also in wild type mice. Mechanistically, based on experiments in HEK293 cells, it was suggested that LRP6 directly binds to APP, leading to an increased cell surface expression of APP favouring non-amyloidogenic processing and decreased Aβ production [117]. Consistent with the above study, chronic administration of small molecule antagonists of canonical Wnt signalling (ICG001 or XAV939) into wild type mice or the J20 AD mouse model also increased amyloidogenic APP processing [155]. Together, these two studies highlight the importance of Wnt signalling for correct APP processing, especially given the fact that Wnt signalling attenuation triggered AD-like amyloidogenic changes (as well as memory deficits) including in wild type mice. Overall, it appears that APP processing and Wnt signalling are engaged in a cross-regulatory network, which deteriorates into a pathologic feedback loop in AD as Wnt signalling is depressed, causing increased amyloidogenic processing, which in turn further depresses Wnt signalling [117]. Consistent with this hypothesis, feedback loop intervention by activation of Wnt signalling with LiCl (a GSK-3β inhibitor), Wnt3a or Wnt7a rescued the neurodegeneration and behavioural impairments induced by Aβ fibrils [156–158], while Aβ sequestration restored canonical Wnt signalling [159] in rat primary neurons as well as in cell lines.

4.2. Wnt Signalling, Tau Pathology, and AD

The second major hallmark of AD is the emergence of tau pathology, otherwise known as tauopathy, in the shape of neurofibrillary tangles (NFTs) within neurons. NFTs are made up of the microtubule associated tau protein in a hyperphosphorylated state (tau usually has a microtubule
stabilising function, which is governed to a large extent by its phosphorylation status that is subject to the action of kinases and phosphatases; [160,161]). Multiple neuronal processes critically depend on proper tau function, including mitochondrial transport, synaptic transmission, and autophagy (reviewed in [162]). Consistent with its important role, mutations in tau-encoding gene MAPT frequently negatively impact microtubule dependent activities especially within neurons, due to the emergence of tau hyperphosphorylation and tauopathy. However, unlike mutations in APP and APP processing genes, mutations in MAPT typically result in in a different type of dementia called frontotemporal dementia (FTD) rather than FAD [163,164]. Nonetheless, tau is inextricably linked to AD as evidenced by the presence of important mechanistic and perhaps synergistic linkages between Aβ and tau but the precise nature of this relationship is complex. For instance, tau pathology may arise independently of cortical Aβ, but equally, tau spreading and cognitive decline are facilitated by the presence of cortical Aβ according to studies on various AD mouse models [165–172].

In addition to phosphorylating and thereby inhibiting β-catenin, GSK-3β is also one of the major kinases responsible for tau phosphorylation [173,174]. Canonical Wnt signalling usually inactivates GSK-3β. However, as Wnt signalling strength is reduced in AD, the activity of GSK-3β is increased, thus contributing to AD-related tau hyperphosphorylation [108,175–179]. As discussed above, Wnt signalling offers pathological opportunities for crosstalk between amyloidogenic APP processing and tau hyperphosphorylation. Accordingly, FAD-associated mutations in the APP processing related gene PSEN1 (known to reduce Wnt signalling; see Section 4.1), increase GSK-3β activation and tau phosphorylation in human brain biopsies, an effect that is likely related to the ability of PS1 to associate with both GSK-3β and tau [180]. Furthermore, studies have confirmed that Aβ treatment activates GSK-3β and increases tau hyperphosphorylation in cultured neurons, acute brain slices, as well as in vivo, which could be mimicked by Dkk1 treatment and blocked with the GSK-3β inhibitor 6-bromo-indirubin-3′-oxime [175,176]. There was also a concomitant decrease in the activity of the tau phosphatase PP2A in rat hippocampal slices, indicating that the usually tightly controlled balance between tau phosphorylation and dephosphorylation is now more generally shifted towards phosphorylation ultimately resulting in tau hyperphosphorylation and tauopathy [175].

Another link between tau and Wnt signalling was recently described, where it was discovered that tau itself can acetylate β-catenin, which increased β-catenin stability and nuclear translocation in HEK cells and rat primary neurons, and further promoted cell survival [181,182]. However, in hippocampal extracts from AD patients, the levels of β-catenin and K49-acetylated β-catenin were increased and further positively correlated with phosphorylation levels of tau at S199 [182]. While decreased β-catenin levels, and hence Wnt signalling, would be more intuitive in the context of AD, it should be pointed out that no distinction between nuclear and total β-catenin levels was made in the above study. Thus, even though a more nuanced evaluation of β-catenin would be required (see also relevant discussion in Section 4.1), β-catenin levels were also increased in PSEN1 related human FAD brains [149,150], indicating that dysregulation of canonical Wnt signalling in AD may manifest via either decreased or increased β-catenin. The authors of [182] speculated whether accumulated tau could help neurons escape acute apoptosis while simultaneously causing chronic neurodegeneration.

4.3. Wnt Signalling, Synapses, and AD

The pathophysiological processes that lead to AD start many years before the diagnosis of AD dementia [6]. During this long preclinical phase, synaptic dysfunction, as determined by 18F fluorodeoxyglucose positron emission tomography or functional magnetic resonance imaging, appears to occur relatively early and before brain atrophy, abnormal cognition, and decreased clinical function become detectable [183]; however, it is important to highlight that the exact timeline with respect to the onset of the individual clinical and pathological hallmarks is still subject to refinement as detection and screening tools are continually improved (e.g., [184]). Nonetheless, it is generally accepted that Aβ pathology emerges before synapse dysfunction. This relationship is not only temporal but also aetiological in nature as Aβ contributes to impaired synaptic function and synapse loss by acting as a
synaptotoxin [185–190]. Given the fact that synapses are crucial for cognitive function and that synapse loss correlates with the degree of cognitive impairment in AD patients, understanding the molecular mechanisms that link Aβ, synapse loss and cognitive decline is thus of utmost importance [191–194].

It has emerged over the years that the endogenous antagonist of canonical Wnt signalling Dkk1 plays a central role in this process. In cultured rat neurons, Aβ oligomer treatment rapidly induced Clu (clusterin) expression and reduced its protein secretion, which resulted in p53-dependent induction of Dkk1 [195]. Of note, DKK1 expression is also evident in neurons of AD patients [119]. This pathway appears to be at least partially tied to synapses since clusterin was demonstrated to accumulate within synapses in human AD brains [196]. As was shown in a study on acute adult mouse hippocampal slices, Dkk1 induction occurs rapidly, where Aβ oligomers induced Dkk1 expression in CA1, CA3 and DG at mRNA and protein levels within 3 h of treatment [121]. Furthermore, in mature rat hippocampal neuron cultures with established synapses, Dkk1 treatment rapidly reduced the number of synaptic sites, both at the pre- and post-synaptic level based on immunocytochemistry and electron microscopy data. This was recapitulated in vivo, where inducible Dkk1 expression resulted in reduced synaptic transmission, impaired LTP, enhanced long term depression (LTD), synapse degeneration and memory deficits in the hippocampus and striatum [111,197]. While LRP5/6 are Fz co-receptors for Wnt ligands, LRP6-Krm1 heterodimers are also the main receptor for Dkk1 and thus Dkk1 acts by removing functional Fz-LRP6 co-receptors for canonical Wnt ligands [45,198–200]. Common LRP6 variants are associated with late onset AD (see Sections 4 and 4.1) while conditional Lrp6 loss of function in neurons elicited synaptic impairment in aged mice and exacerbated amyloid pathology in the APP/PS1 AD mouse model [115,117]. Equally, silencing Krm1 with miR-431 prevented Aβ-related synapse loss in cultured neurons obtained from the 3xTg AD mouse model [200]. Thus, Dkk1-mediated synapse disassembly is mediated by the attenuation of canonical Wnt signalling [111,136,195,201].

Crucially, synapse loss as a response to Dkk1 exposure was reversible in vitro and in vivo, where synapse number and function was restored following removal of Dkk1 by washout or by cessation of Dkk1 over-expression [111,121], indicating that the normalisation of Wnt signalling may enable restoration of synapses in AD patients. In fact, activation of both canonical and non-canonical pathways were proven to counteract the synaptotoxic effects of Aβ: a potentiator of Wnt3a-mediated canonical signalling, WASP-1, as well as the non-canonical ligand Wnt5a both rescued hippocampal synaptic impairments triggered by Aβ oligomers [154,190]. However, there is now also mounting evidence suggesting that activation of non-canonical PCP signalling may exacerbate Aβ oligomer synaptotoxicity. Indeed, Aβ-mediated Dkk1 induction activated the Wnt/PCP pathway via Fz-Vangl2 co-receptors and the resulting modulation of cytoskeletal dynamics contributed to synapse loss, which could be reversed in rat primary cortical neurons by blocking this pathway using inhibitors of the actin cytoskeleton regulatory protein ROCK [111,136,195,201]. The mechanism of action of Dkk1 may thus involve both the reduction of canonical Wnt signalling as well as the activation of the Wnt/PCP pathway, both of which may result in the destabilisation and eventual loss of synapses [136,201]. Adding a further layer of complexity, APP physically interacted with LRP6 and Vangl2 and co-activated canonical and Wnt/PCP signalling respectively in cell line reporter assays. However, over-expression of an FAD-related APPSwe variant was less able to co-activate canonical signalling but more effective at co-activating the Wnt/PCP pathway compared with wild type APP, supporting the idea of a pathologic shift from canonical to non-canonical signalling in AD that is detrimental for synapses [136].

4.4. Wnt Signalling, Microglia/Neuroinflammation, and AD

Microglia are the resident immune cells of the CNS [202]. Using motile processes, they constantly survey their surroundings in search of pathogens and cellular debris that need to be cleared, thus maintaining a healthy CNS environment in a tissue where extensive immune reactions, akin to what is observed in the periphery, would be detrimental. Microglia also play an important role in refining neuronal circuits both during development and in adulthood by removing or stimulating the formation of synapses as well as by removing neuronal precursors [203]. In AD however, microglia
contribute profoundly to pathology. The inherent link between microglia and AD is evidenced by the fact that variants of an increasing number of genes expressed in microglia are associated with an increased or decreased risk of AD, such as variants of TREM2 and CD33 encoding microglial immune receptors [133,204]. The AD brain mounts an inflammatory response involving microglia (as well as astrocytes) [205]. These usually respond to CNS insults by proliferating as well as by altering their gene expression, morphology and secretomes as they transition from a senescent/surveying into an activated state, while extending their processes and migrating towards the lesion site where they initiate an innate immune response (reviewed extensively in [206]). Several RNASeq studies focusing on a range of AD mouse models have identified transcriptionally distinct subsets of microglia that emerge in and are uniquely associated with AD and have collectively been termed ‘Disease Associated Microglia’ (DAM) or ‘Activated Response Microglia’ (ARM) [122,207,208].

The transition from surveying microglia to DAM/ARM cells appears to require TREM2 [207]. TREM2 in return signals intracellularly to affect the expression of numerous genes. In keeping, when Trem2 was knocked out in the PS2APP AD mouse model, the transcript levels of 144 genes were reduced, compared to only 7 genes that were upregulated [209]. As the authors of this study stated, many of these downregulated genes were upregulated in microglia in PS2APP mice in the first place compared to wild type controls when both groups had intact Trem2 [122,209,210]. This suggests that the microglial AD response involves the Trem2-regulated upregulation of numerous genes. This group of genes was enriched for positive regulators of canonical Wnt signalling according to gene ontological analysis (Fz9, Sult2, Bambi, Ptk7, Aspin, and Dkk2) [209]. Three further Trem2-regulated genes—Wif1, Ctnna3 and Dkk2—can be categorised as negative regulators of Wnt signalling (note that Dkk2 can be a context-dependent activator and inhibitor of canonical Wnt signalling [47]) and their biological function in this context is most likely related to negative feedback [209].

Interestingly, the existence of crosstalk between TREM2 and canonical Wnt signalling in microglia had already been discovered in an earlier mouse study [134], where it was demonstrated that the proliferative response of activated microglia not only requires TREM2 but also crosstalks with canonical Wnt signalling [134]; although not explicitly addressed within this study, it appears plausible that the same would apply to microglial proliferation in human AD brain. Trem2−/− knockout mice exhibited impaired kainic acid-induced microglial proliferation and survival, while microglial apoptosis was increased. Simultaneously, β-catenin and inactive GSK-3β protein levels were reduced. Based on these data it was proposed that TREM2 usually signals intracellularly to inactivate GSK-3β via phosphorylation by PI3K/AKT, leading to β-catenin stabilisation and transcription of pro-survival and anti-apoptotic genes. Obviously, GSK-3β and AKT are important nodal points that provide links to canonical Wnt signalling. In keeping, direct and indirect GSK-3β activation with LiCl and Wnt3a respectively rescued the microglial phenotype in Trem2−/− mice.

However, the exact function of the microglial response in the AD brain is multifaceted and whether it is regarded as beneficial or detrimental appears to be context dependent. This dichotomy can be exemplified by the way microglia respond to Aβ. It is widely established that activated microglia accumulate around Aβ plaques using the Aβ cell surface receptors SCARA1, CD36, CD14, α6β1 integrin, CD47, TLR2/4/6/9, TREM2 and that they can phagocytose Aβ (note the link between Wnt signalling and Aβ as well as TREM2 already described in earlier sections) [211–214]. This contributes to the notion that microglia play a protective/beneficial role, perhaps at certain stages of AD, even though they ultimately fail to efficiently clear Aβ in AD. Microglia form a physical barrier around the fibrillar plaque centre while limiting the halo-like leakage of the more neurotoxic oligomeric Aβ from these centres [215] (note however, that disparate findings have been published where plaque formation/maintenance was affected by microglia ablation in some cases but not in others [209,216–218]). In addition, microglia also orchestrate plasminogen and metalloproteinase proteolytic activities to act on Aβ [219]. These beneficial roles are complemented by the clearing of AD-related cellular debris and dead cells.

By contrast, microglia also exert detrimental effects in AD that exacerbate disease. Cytokines and chemokines secreted by activated microglia (and astrocytes) are important for the chemoattraction and
response towards Aβ and cellular debris, likely helping to mitigate disease at early stages. However, chronically activated microglia continually secrete pro-inflammatory cytokines (e.g., TNFα, IL-6, IL-1α, and GM-CSF), which is an important, albeit harmful, aspect of neuroinflammation evident in the AD brain, leading to a sustained pro-inflammatory environment as well as suppression of neuronal activity and injury of bystander neurons [220]. Furthermore, microglia increasingly engage in synaptic pruning using the complement system, thus contributing to the deterioration of neuronal circuits [221,222]. All these activities are likely exacerbated due to the microglial proliferative response regulated by canonical Wnt signalling. Thus, whether microglia are friend or foe in the fight against AD may thus very well be a context-dependent matter.

The molecular mechanism underlying the microglial friend/foe dichotomy may be centred around the interaction between Wnt and Toll like receptor (TLR) signalling pathways [223,224]. However, this relationship is presently still ill-defined. Key molecular components of these pathways have been linked in a limited number of AD studies focusing on microglia, but a lot of insight into detailed cross-talk mechanisms were initially derived from peripheral immune cell or cancer responses. TLRs belong to the family of pattern recognition receptors that detect pathogenic or damage-derived molecules including Aβ, and many TLRs are expressed by immune cells such as microglia, although they are also expressed in various other CNS cells. In AD, Aβ is recognised by TLR2 and TLR4 on microglia [225,226] and in the classical TLR pathway, ligand binding culminates in the expression of pro-inflammatory genes including genes encoding IL-1β, IL-6, IL-12, TNF-α, iNOS, and COX2, driven by the transcription factor NF-κB [214]. Importantly, canonical Wnt signalling, via both GSK-3β and β-catenin, can exert both anti- and pro-inflammatory activity in non-neural models, respectively preventing and stimulating the expression of pro-inflammatory NF-κB target genes, suggesting that Wnt signalling modulates TLR signalling [227–229]. These divergent findings imply a context-dependent cross-talk that could explain the transition towards the pro-inflammatory state exhibited by chronically activated AD microglia. This might occur in a concerted fashion with TREM2-mediated, induced proliferation of microglia that is supported by canonical Wnt signalling [134]. In keeping, a recent study demonstrated that Wnt3a treatment ameliorated toxic microglial responses in an ischemic stroke mouse model by reducing peri-infarct microgliosis and decreasing expression of iNos and TNF-α, two known NF-κB target genes [230]. Another recent study showed in a developmental encephalopathy mouse model that the canonical Wnt agonist L803mts targeted at microglia negatively modulated microglial activation, manifesting as a reduction in iNos and Cox2 mRNA levels in microglia [229]. The possibilities for Wnt/TLR cross-talk in microglia are ample and bi-directional (Wnt signalling can affect TLR signalling and vice versa) extending well beyond just GSK-3β and β-catenin although direct evidence for this cross-talk in microglia is only beginning to be discovered. The reader is referred to a previously published review for further reference [224].

4.5. Wnt Signalling, the Blood Brain Barrier, and AD

The blood brain barrier (BBB) is a restrictive barrier between the general circulation and the CNS serving to maintain correct brain parenchyma and cerebrospinal fluid composition by controlling influx/efflux of molecules and cells into/out of the CNS and to protect the CNS from peripheral insults [231]. In the wider context, the BBB is an essential component of the neurovascular unit that is further comprised of neurons, glia (variously including astrocytes, microglia, and oligodendrocytes), which form a tightly integrated system that ensures normal brain function. The BBB consists of endothelial cells (ECs) of the CNS microvasculature, as well as a group of perivascular components comprising astrocytes, pericytes, and the extracellular matrix. Much of the barrier function stems from two different protein structures between ECs. Tight junctions are comprised of Occludin, Claudins, and other junctional adhesion molecules assembled into a barrier forming zip-locked structure that restricts diffusion of solutes from the blood into the CNS [232]. Adherens junctions contain transmembrane cell adhesion proteins of the cadherin family (mostly VE-cadherin) that are responsible for cell-cell adhesion between CNS ECs. Intracellularly, adherens junctions are linked
to the actin cytoskeleton via catenins (α-, β-, and γ-catenin), which regulate the adhesive properties of VE-cadherin (note that β-catenin can have dual roles in cell adhesion and Wnt signalling in this context), actin bundling that is linked to barrier permeability, as well as cell signalling processes [231]. To enable the passage of otherwise non-permeant solutes into and out of the CNS, the BBB is equipped with transporters such as GLUT-1 (supplying the brain with its primary energy source glucose) and P-gp (broad spectrum efflux transporter restricting solute access to the CNS or expelling solutes such as Aβ into the blood) [233,234].

Given these important roles, BBB dysfunction can have detrimental consequences on the CNS. Indeed, BBB disruptions are a key feature of AD, bringing about increased BBB permeability leading to CNS microbleeding and perivascular infiltration of neurotoxic blood-derived products, cellular infiltration and degeneration of pericytes and ECs, as well as impaired GLUT-1 and P-gp function, which negatively impact glucose homeostasis and Aβ clearance (reviewed in [235,236]). These defects may be in part linked to impaired Wnt signalling at the BBB; this is in light of the well-established role that this signalling pathway plays in CNS vasculature/BBB development and maintenance: Wnt1, Wnt3a, Wnt4, Wnt7a, Wnt7b, and Norrin have been identified as neuroepithelium derived Wnt ligands targeting the vascular endothelium that expresses cognate receptors Fz4, 6, and 8 as well as the co-receptors Lrp5/6, Gpr124, and Reck to stimulate vascular development and barrier formation specifically in the CNS via the canonical pathway [51,237–243]. Furthermore, canonical Wnt signalling, through transcriptional transactivation or protein-protein interactions, regulates target genes or proteins relevant for CNS vasculature/BBB function, including Claudin-3, VE-cadherin, GLUT-1, and P-gp in brain endothelial cells from zebrafish, rodents and humans [237,244–247]; this may occur in coordination with non-canonical Wnt pathways via RhoA and RhoAK-mediated inhibition of GSK-3β activation [248]. The fact that global knockout or endothelial cell-restricted conditional knockout of canonical Wnt components such as Wnt7a, Fz4, Norrin or Ctnnb1 (β-catenin) in adult mice reduced claudin-1, -3, -5, and GLUT-1 protein levels accompanied in some cases by seizures, neuronal injury, haemorrhages, and inflammation demonstrates that canonical Wnt signalling is also important for CNS vasculature/BBB maintenance [249,250].

As mentioned earlier, the AD brain suffers from increased Aβ burden and impaired Wnt signalling as part of a pathological negative feedback loop (see Section 4.1). This downward spiral is further exacerbated by the fact that reduced canonical Wnt signalling strength, as brought about by Dkk1 and Aβ treatment or in transgenic mice expressing constitutively active Gsk3b (S9A), was shown to decrease MDR1 (P-gp) expression in brain endothelial cells in vitro and in vivo as well as in the hippocampus of AD patients [246,251–253]. In absence of P-gp-mediated Aβ expulsion into the general circulation, Aβ retention within the brain is increased [254] (Note that Aβ removal from brain to general circulation involves the crossing of endothelial cells, first on the abluminal (brain side) plasma membrane, aided by LRP1, then on the luminal (blood side) plasma membrane, mediated by P-gp). This is likely exacerbated by the fact that expression of the receptor for advanced glycation end-products (RAGE), believed to mediate Aβ uptake from the blood into the brain, is increased [255]. Conversely, restoring Mdr1 expression in the hAPP AD mouse model, significantly reduced brain Aβ levels [256].

Reduced Wnt signalling in AD may have direct consequences on energy metabolism in the brain. Canonical Wnt signalling via Wnt7a was shown to boost expression of Slc2a1 (GLUT-1) in primary mouse brain endothelial cells [237]. With glucose being the primary energy source in the brain, reduced endothelial cell GLUT-1 levels, as observed in the AD brain [257], can lead to insufficient glucose tissue uptake. Indeed, in the Slc2a1 deficient APPsw/0 AD mouse model there is exacerbated AD pathology exemplified by accelerated microvascular degeneration, blood flow reductions, BBB breakdown, accelerated Aβ pathology and reduced clearance, attenuated neuronal activity, and more severe neurodegeneration [258].

Together, these lines of evidence demonstrate the importance of Wnt signalling in BBB development and maintenance. Consequently, dysregulation of Wnt pathways could contribute profoundly to the vascular and, by association, the neuro-glial, pathobiology evident in AD.
4.6. Are the Wnt Changes Seen in AD Simply an Exacerbated Consequence of Ageing?

The identification of genetic risk factors in the previous sections notwithstanding, age still remains the biggest risk factor for AD [112,113,259]. Indeed, a recent review has pointed out that a number of hallmarks of ageing are either also found in AD patients or can be linked to exacerbated disease in experimental models and patients [259]; these include increased DNA damage/reduced DNA repair, telomere instability, epigenetic changes, compromised mitophagy, increased senescence, stem cell depletion, increased neuroinflammation, and impaired brain metabolism. Could the same also apply to dysregulated Wnt signalling?

We, and others in the past (e.g., [11]) have discussed that dysregulated Wnt signalling is indeed another commonality between ageing and AD, manifesting as an overall decrease in the Wnt signalling tone caused by decreased expression of Wnt ligands and receptors as well as increased expression and/or activity of negative regulators of Wnt signalling such as GSK-3β ([108]; see also Section 3.3). However, while increased expression of the endogenous canonical Wnt signalling antagonist Dkk1 has intriguingly been identified both in old age and AD mouse models, DKK1 upregulation has so far only been identified in AD patients, but not in normal ageing [108,119]. Comparisons between normal ageing brain and AD brain are presently difficult due to the relative scarcity of published literature on the former at the time of writing. The plethora of single cell and single nuclear sequencing studies on AD/control brain that are now emerging may shed further light on this question [260].

In spite of some apparent overall commonalities with respect to dysregulated Wnt signalling in normal ageing and AD, there is presently a gap in our understanding of the underlying molecular mechanisms and whether they are similar or conserved. It can be speculated whether the other commonalities between normal ageing and AD mentioned at the beginning of this section, might result in impaired Wnt signalling (e.g., epigenetic changes, neuroinflammation etc.). Conversely, although still far from a complete mechanistic description, links between dysregulated Wnt signalling and AD have previously been described (e.g., the connection between Aβ, CLU and upregulated DKK1 expression [195]; see also previous segments in Section 4). It is conceivable that subthreshold pathological changes, including increased amyloidogenic APP processing, could destabilise normal Wnt signalling both in AD and normal ageing. While these changes evolve during the long prodromal AD stage, they could remain below the pathological threshold in normal ageing. This would be indicative of a quantitative difference in the relative amount of Wnt signalling dysregulation between these two scenarios. However, in absence of supporting evidence, qualitatively different molecular mechanisms are presently equally conceivable. The fact that the molecular fluid biomarker signature between normal ageing and AD is different (e.g., [184]) would support qualitative differences between the two. In this context, Wnt signalling components are yet to be firmly established as fluid biomarkers for AD, although recent studies have suggested serum DKK1 as a predictor of deteriorating disease both in AD and acute ischemic stroke patients [261,262].

5. Looking Forward: Does Wnt Signalling Offer Opportunities for New Therapies for AD?

Normalising dysfunctional Wnt signalling in the AD brain may represent an opportunity for novel therapeutic approaches. As has been discussed, the data points towards a decreased Wnt signalling tone and therefore a therapeutic intervention would necessitate restored signalling. This flags a potential safety concern given the extensive data on the role of dysregulated (generally an increased) Wnt signalling in cancer [263]. Similarly, Wnt signalling pathways exhibit a high degree of complexity with a plethora of cross-talk points with other signalling pathways and cellular processes, potentially making it difficult to target this signalling pathway in a controlled way without causing unwanted side effects (although some therapeutic success in experimental models has been achieved as summarised in Table 1). Thus from this perspective there is a need to proceed with caution. Generic upregulation of Wnt signalling (for instance increasing canonical Wnt signalling through use of a small molecule GSK-3β inhibitor) is unlikely to be a viable approach for treating a chronic disease. One way to proceed may be to identify ways of selectively targeting Wnt signalling (as opposed to generically
increasing signalling). This could be achieved for instance by (i) targeting specific components of Wnt signalling that are dysregulated in the disease, or (ii) by targeting signalling pathway components that are selectively expressed in specific cell types. Additionally, while as has been discussed (Section 4.1) there is an evidence base to support the modulation of APP processing/β-amyloid production by Wnt signalling, from a therapeutic perspective it can be argued that there are more direct ways to target the β-amyloid cascade [264], and the lack of success in clinical trials to date, has led to a general deprioritisation of β-amyloid as a target [265]. As such, this will not be considered further.

The secreted Wnt antagonist DKK1 is upregulated in AD and directly implicated in synapse dysfunction and loss (see Section 4.3); as such it is a potential therapeutic target. One approach that has been investigated in the context of another chronic disease, osteoporosis, is the use of an anti-DKK1 monoclonal antibody to neutralise its inhibitory activity; indeed, this has been shown to be efficacious in animal models [266]. The obvious caveat for the application of this approach to AD is our current inability to deliver large therapeutic molecules such as antibodies across the BBB. A second approach would be to develop a small molecule drug that could inhibit the binding of DKK1 to its cell surface co-receptor LRP6. However, developing small molecule drugs that inhibit the interaction of large proteins has generally proven to be very challenging [267]. The third approach may be to employ antisense oligonucleotide (ASO) technology to target DKK1. ASOs are in development for a number of disorders including neurodegenerative diseases such as Huntington’s disease [268].

Table 1. Summary of Wnt targeting approaches in the published literature, where an increase of or restoration to normal signalling levels was attempted to treat neurological disorders (predominantly AD) in experimental models. ‘↑’ and ‘↓’ indicate increases or decreases in a measured experimental outcome.

| Reference                  | Type of Manipulation | Experimental Model                      | Experimental Outcome                                                                 |
|----------------------------|----------------------|----------------------------------------|---------------------------------------------------------------------------------------|
| Parr et al., 2014 [153]    | Wnt3a, β-catenin over-expression | N2A5 sw murine neuroblastoma cell line | ↓ Aβ levels, ↓ BACE1 levels                                                            |
| Alvarez et al., 2004 [157] | Wnt3a treatment       | Rat primary hippocampal neurons (treated with Aβ) | ↑ neuronal survival, ↓ GSK-3β phosphorylation, ↓ tau phosphorylation, restored cytosolic β-catenin levels, ↑ increased expression of Wnt target gene engrailed-1 |
| Zhang et al., 2019 [230]   | Wnt3a                 | Ischemic stroke (mMCAO) mouse model (in vivo) | ↓ infarct volume, ↓ neurologic deficits, ↓ peri-infarct apoptosis, ↓ microglial activation, ↓ pro-inflammatory cytokine secretion (iNos, TNFα), ↓ astrogliosis |
| Zheng et al., 2017 [134]   | Wnt3a, LiCl, TDZD-8    | Trem2-/- mouse primary microglia        | ↑ β-catenin levels, ↑ microglial survival, ↑ microglial proliferation                  |
| Cerpa et al., 2010 [196]   | Wnt5a                 | Rat primary hippocampal neurons and hippocampal slices (treated with Aβ) | ↑ fEPSP & (AMPAR, NMDAR) EPSC amplitudes, ↑ post-synaptic PSD-95 clusters              |
| Marzo et al., 2016 [111]   | 6-BIO (GSK-3β inhibitor) | Rat primary hippocampal neurons (treated with Dkk1) | ↑ excitatory synapse numbers                                                             |
| Salcedo-Tello et al., 2014 | 6-BIO                 | Rat hippocampal slices at 3 months (young) and 18–20 months (aged) | ↓ GSK-3β activity, ↑ β-catenin levels                                                  |
| Quintanilla et al., 2005   | 17β-estradiol, Trolox (antioxidant) | Rat primary hippocampal neurons (treated with Aβ) | ↑ neuronal survival, ↓ endoperoxide production, ↓ GSK-3β activity, ↑ cytoplasmic β-catenin, ↑ Wnt5a & Wnt7a levels (17β-estradiol), ↓ Ca²⁺ mediated toxicity |
|                           | Wnt7a                 |                                         | protection against Ca²⁺ mediated toxicity                                                |
Table 1. Cont.

| Reference                        | Type of Manipulation                | Experimental Model                                           | Experimental Outcome                                                                 |
|----------------------------------|------------------------------------|--------------------------------------------------------------|---------------------------------------------------------------------------------------|
| Magdesian et al., 2008 [159]     | cSP5 (Aβ-binding peptide)          | N2Asmurine neuroblastoma cell line (differentiated)         | ↑ total & nuclear β-catenin                                                           |
| Elliott et al., 2018 [136]       | Fasudil (Rock inhibitor)           | Rat primary cortical neurons (treated with Dkk1)             | Protection against Dkk1-mediated dendritic spine loss, and Aβ production              |
| Vargas et al., 2014 [187]        | FOXY-5 (peptide)                   | APP/PS1 mouse model (in vivo & hippocampal slices)           | Rescued memory retention, ↑ fEPSP; rescued LTP deficit                               |
| Van Steenwinckel et al., 2019 [229] | L803mts (Wnt/β-catenin activator; targeted to microglia by 3DNA) | Mouse developmental encephalopathy model (IL-1β-mediated; in vivo) | ↓ microglial expression of pro-inflammatory cytokines (iNos, Cox2), restored myelination, restored short- & long-term memory retention |
| De Ferrari et al., 2003 [156]    | LiCl (GSK-3β inhibitor), Wnt3a, PMA (PKC activator) | Rat primary hippocampal neurons (treated with Aβ)            | ↑ neuronal survival, restored cytosolic β-catenin levels                             |
| Scali et al., 2006 [176]         | LiCl                               | Rats (3 months) injected with recombinant human DKK1 (in vivo) | ↓ DKK1-stimulated GSK-3β activity, protects against DKK1-mediated neurodegeneration, tau hyperphosphorylation, and astrogliosis |
| Ross et al., 2018 [200]          | miRNA-431 (silences Dkk1 receptor Krm1) | Mouse primary cortico-hippocampal neurons from 3xTg AD mouse model (in vivo) | ↓ Aβ- and Dkk1-mediated synapse loss, ↓ neurite degeneration |
| Vargas et al., 2015 [154]        | WASP-1 small molecule              | Rat primary hippocampal neurons                              | ↑ synaptic transmission, rescued hippocampal LTP impairments due to Aβ                |
| Y27632 (Rock inhibitor)          | Y27632, Fasudil                    | Rat primary cortical neurons (treated with Dkk1)             | Protection against Aβ- or Dkk1-mediated spine density reduction, ↑ PSD-95 puncta, ↑ GluA1 puncta |
| Sellers el al., 2018 [201]       | Fasudil                            | Rats (treated with Aβ; in vivo)                              | Protection against Aβ-driven cognitive impairment, ↑ performance in NOR task          |
|                                 | LiCl                               | Trem2-/- mice (in vivo)                                     | ↑ microglia cell # in hippocampus and cortex, ↑ β-catenin, ↓ GSK-3β activation, ↑ cell cycle markers (c-Myc, Cyclin D1) |

In addition to considering the therapeutic modality, there are some other points to consider. While DKK1 expression is increased in AD brain (Section 4.3), the stage of the disease when this occurs has not been defined. Such understanding is essential to inform when to intervene with a DKK1 based therapeutic. Similarly, can DKK1 be detected in cerebrospinal fluid (CSF)? Further establishing DKK1 CSF or serum biomarkers [261,262] would be valuable in understanding where DKK1 fits in the pathophysiology of AD, as well as being a target biomarker for evaluating DKK1 therapeutics. It would also have the potential to enable precision medicine by identifying those patients who have upregulated DKK1.
Microglia and neuroinflammation have become an area of significant activity in the last few years, including as potential targets for therapeutic intervention in AD. A particular focus has been the TREM2 signalling pathway, where there are efforts being explored to stimulate this pathway and as a therapeutic approach for AD. We have discussed above (Section 4.4) that there are a number of studies describing an interaction between TREM2 and Wnt signalling in microglia. While this offers the potential for modulating TREM2 signalling via Wnt, it is premature from a therapeutic perspective. Our understanding of the interaction between these signalling pathways is currently limited. Importantly, there is currently no data as to whether this occurs in human microglia, and whether there is dysfunction or dysregulation in AD that could contribute to the pathology of the disease.

In Section 4.5 the importance of the Wnt signalling system in maintaining a functional BBB was discussed, as was the dysfunction of the BBB in AD. While the loss of Wnt signalling in brain endothelial cells contributes to BBB breakdown in disorders such as multiple sclerosis, hemorrhagic stroke, and traumatic brain injury, no such evidence is currently available for AD, either in animal models or human disease. This is clearly a gap in our knowledge, and a gap in terms of supporting the notion of modulating BBB Wnt signalling as a therapeutic approach for AD. Nevertheless, given the decreased expression in AD brain of Wnt regulated genes essential for normal BBB function such as endothelial cell tight junction proteins and transporters, it is a plausible hypothesis that increasing Wnt signalling tone at brain endothelial cells has the potential to restore expression and therefore be of therapeutic benefit. Indeed, the Wnt co-receptor Gpr124, through which endothelial cell signalling is mediated is selectively expressed in brain endothelial cells and so in principle could be targeted to generate agonists or positive allosteric modulators, as has been done for other members of the GPCR gene super-family.

In summary, while our understanding of the dysregulation of Wnt signalling in AD has increased, there remain significant gaps, particularly a clear understanding of the dysfunction in the human disease, and how this links to the pathophysiology. This deeper knowledge will be essential in informing rational approaches to developing new therapies that could become part of our armoury for treating AD in the future. However, the success of future Wnt-therapies will inevitably require early intervention and hence early detection at pre-clinical stages. Decreased Wnt signalling strength is evident in the ageing brain and likely further declines in the transition to pathological ageing, mild cognitive impairment, and AD. Future research efforts should thus also focus on the development of early detection methods for reduced Wnt signalling strength, as proposed in this section for DKK1, in potential future LOAD patients at prodromal stages.

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References

1. Albert, M.S.; DeKosky, S.T.; Dickson, D.; Dubois, B.; Feldman, H.H.; Fox, N.C.; Gamst, A.; Holtzman, D.M.; Jagust, W.J.; Petersen, R.C.; et al. The diagnosis of mild cognitive impairment due to Alzheimer’s disease: Recommendations from the National Institute on Aging-Alzheimer’s Association workgroups on diagnostic guidelines for Alzheimer’s disease. Alzheimer’s Dement. 2011, 7, 270–279. [CrossRef]

2. McKhann, G.M.; Knopman, D.S.; Chertkow, H.; Hyman, B.T.; Jack, C.R., Jr.; Kawas, C.H.; Klunk, W.E.; Koroshetz, W.J.; Marly, J.J.; Mayeux, R.; et al. The diagnosis of dementia due to Alzheimer’s disease: Recommendations from the National Institute on Aging-Alzheimer’s Association workgroups on diagnostic guidelines for Alzheimer’s disease. Alzheimer’s Dement. 2011, 7, 263–269. [CrossRef] [PubMed]
3. Alzheimer’s Association. 2019 Alzheimer’s disease facts and figures. *Alzheimer’s Dement.* 2019, 15, 321–387. [CrossRef]
4. Selkoe, D.J.; Hardy, J. The amyloid hypothesis of Alzheimer’s disease at 25 years. *EMBO Mol. Med.* 2016, 8, 595–608. [CrossRef]
5. Fish, P.V.; Steadman, D.; Bayle, E.D.; Whiting, P. New approaches for the treatment of Alzheimer’s disease. *Bioorg. Med. Chem. Lett.* 2019, 29, 125–133. [CrossRef]
6. Long, J.M.; Holtzman, D.M. Alzheimer Disease: An Update on Pathobiology and Treatment Strategies. *Cell 2019*, 179, 312–339. [CrossRef] [PubMed]
7. Cao, J.; Hou, J.; Ping, J.; Cai, D. Advances in developing novel therapeutic strategies for Alzheimer’s disease. *Mol. Neurodegener.* 2018, 13, 1–20. [CrossRef]
8. Vermunt, L.; Sikkes, S.A.; Hout, A.V.D.; Handels, R.; Bos, I.; Van Der Flier, W.M.; Kern, S.; Ousset, P.-J.; Maruff, P.; Skoog, I.; et al. Duration of preclinical, prodromal, and dementia stages of Alzheimer’s disease in relation to age, sex, and APOE genotype. *Alzheimer’s Dement.* 2019, 15, 888–898. [CrossRef]
9. Jagust, W.; Jessen, F.; Karlawish, J.; et al. NIA-AA Research Framework: Toward a biological definition of Alzheimer’s disease. *Alzheimer’s Dement.* 2018, 14, 535–562. [CrossRef] [PubMed]
10. Sayas, C.L. Tau-based therapies for Alzheimer’s disease: Promising novel neuroprotective approaches. In *Neuroprotection in Autism, Schizophrenia and Alzheimer’s Disease*; Elsevier BV: Amsterdam, The Netherlands, 2020; pp. 245–272.
11. Palomer, E.; Buechler, J.; Salinas, P.C. Wnt Signaling Deregulation in the Aging and Alzheimer’s Brain. *Front. Cell. Neurosci.* 2019, 13, 227. [CrossRef]
12. Inestrosa, N.C.; Varela-Nallar, L. Wnt signaling in the nervous system and in Alzheimer’s disease. *J. Mol. Cell Biol.* 2014, 6, 64–74. [CrossRef] [PubMed]
13. Willert, K.; Brown, J.D.; Danenberg, E.; Duncan, A.W.; Weissman, I.L.; Reya, T.; Yates, J.R.; Nusse, R. Wnt proteins are lipid-modified and can act as stem cell growth factors. *Nat. Cell Biol.* 2003, 423, 448–452. [CrossRef] [PubMed]
14. Logan, C.Y.; Nusse, R. The Wnt signaling pathway in development and disease. *Annu. Rev. Cell Dev. Biol.* 2004, 20, 781–810. [CrossRef] [PubMed]
15. Clevers, H.; Nusse, R. Wnt/beta-catenin signaling and disease. *Cell 2012.* [CrossRef] [PubMed]
16. Schulte, G.; Bryja, V. The Frizzled family of unconventional G-protein-coupled receptors. *Trends Pharmacol. Sci.* 2007, 28, 518–525. [CrossRef]
17. Macdonald, B.T.; He, X. Frizzled and LRP5/6 Receptors for Wnt/β-Catenin Signaling. *Cold Spring Harb. Perspect. Biol.* 2012, 4, a007880. [CrossRef]
18. Fradkin, L.G.; Dura, J.-M.; Noordermeer, J.N. Ryks: New partners for Wnts in the developing and regenerating nervous system. *Trends Neurosci.* 2010, 33, 84–92. [CrossRef]
19. Green, J.L.; Kunz, S.G.; Sternberg, P.W. Ror receptor tyrosine kinases: Orphans no more. *Trends Cell Biol.* 2008, 18, 536–544. [CrossRef]
20. Bhanot, P.; Brink, M.; Samos, C.H.; Hsieh, J.-C.; Wang, Y.; Macke, J.P.; Andrew, D.; Nathans, J.; Nusse, R. A new member of the frizzled family from Drosophila functions as a Wingless receptor. *Nat. Cell Biol.* 1996, 382, 225–230. [CrossRef]
21. Janda, C.Y.; Waghray, D.; Levin, A.M.; Thomas, C.; Garcia, K.C. Structural Basis of Wnt Recognition by Frizzled. *Science 2012*, 337, 59–64. [CrossRef]
22. Dann, C.E.; Hsieh, J.-C.; Ratnner, A.; Sharma, D.; Nathans, J.; Leahy, D.J. Insights into Wnt binding and signalling from the structures of two Frizzled cysteine-rich domains. *Nat. Cell Biol.* 2001, 412, 86–90. [CrossRef] [PubMed]
23. Kikuchi, A.; Yamamoto, H.; Sato, A.; Matsumoto, S. New Insights into the Mechanism of Wnt Signaling Pathway Activation. *Int. Rev. Cell Mol. Biol.* 2011, 291, 21–71. [CrossRef] [PubMed]
24. Kikuchi, A.; Yamamoto, H.; Sato, A. Selective activation mechanisms of Wnt signaling pathways. *Trends Cell Biol.* 2009, 19, 119–129. [CrossRef] [PubMed]
25. MacDonald, B.T.; Tamai, K.; He, X. Wnt/β-Catenin Signaling: Components, Mechanisms, and Diseases. *Dev. Cell* 2009, 17, 9–26. [CrossRef]
26. Liu, C.; Li, Y.; Semenov, M.; Han, C.; Baeg, G.-H.; Tan, Y.; Zhang, Z.; Lin, X.; He, X. Control of β-Catenin Phosphorylation/Degradation by a Dual-Kinase Mechanism. *Cell 2002*, 108, 837–847. [CrossRef]
27. Huber, O.; Korn, R.; McLaughlin, J.; Ohsugi, M.; Hermann, B.G.; Kemler, R. Nuclear localization of β-catenin by interaction with transcription factor LEF-1. *Mech. Dev.* 1996, 59, 3–10. [CrossRef]

28. Molenaar, M.; Van De Weering, M.; Oosterwegel, M.; Peterson-Maduro, J.; Godsave, S.; Korinek, V.; Roose, J.; Destrée, O.; Clevers, H. Xclrf-3 Transcription Factor Mediates β-Catenin-Induced Axis Formation in Xenopus Embryos. *Cell* 1996, 86, 391–399. [CrossRef]

29. Vlad, A.; Röhrs, S.; Klein-Hitpass, L.; Müller, O. The first five years of the Wnt targetome. *Cell. Signal.* 2008, 20, 795–802. [CrossRef]

30. Macheda, M.L.; Sun, W.W.; Kugathasan, K.; Hogan, B.M.; Bower, N.I.; Halford, M.M.; Zhang, Y.F.; Jacques, B.E.; Lieschke, G.J.; Dabdoub, A.; et al. The Wnt Receptor Ryk Plays a Role in Mammalian Planar Cell Polarity Signaling. *J. Biol. Chem.* 2012, 287, 29312–29323. [CrossRef]

31. Nishita, M.; Itsukushima, S.; Nomachi, A.; Endo, M.; Wang, Z.; Inaba, D.; Qiao, S.; Takada, S.; Kikuchi, A.; Minami, Y. Ror2/FRizzled Complex Mediates Wnt5a-Induced AP-1 Activation by Regulating Dishevelled Polymerization. *Mol. Cell. Biol.* 2010, 30, 3610–3619. [CrossRef]

32. Green, J.; Nusse, R.; Van Amerongen, R. The Role of Ryk and Ror Receptor Tyrosine Kinases in Wnt Signal Transduction. *Cold Spring Harb. Perspect. Biol.* 2014, 6, a009175. [CrossRef] [PubMed]

33. Strutt, D.; Weber, U.; Mlodzik, M. The role of RhoA in tissue polarity and Frizzled signalling. *Nat. Cell Biol.* 1997, 387, 292–295. [CrossRef] [PubMed]

34. Fanto, M.; Weber, U.; Strutt, D.; Mlodzik, M. Nuclear signaling by Rac and Rho GTPases is required in the establishment of epithelial planar polarity in the Drosophila eye. *Curr. Biol.* 2000, 10, 979–988. [CrossRef]

35. Boutros, M.; Paricio, N.; Strutt, D.I.; Mlodzik, M. Dishevelled Activates JNK and Discriminates between JNK Pathways in Planar Polarity and wingless Signaling. *Cell* 1998, 94, 109–118. [CrossRef]

36. Winter, C.G.; Wang, B.; Ballew, A.; Royou, A.; Karess, R.; Axelrod, J.D.; Luo, L. Drosophila Rho-Associated Kinase (Drok) Links Frizzled-Mediated Planar Cell Polarity Signaling to the Actin Cytoskeleton. *Cell* 2001, 105, 81–91. [CrossRef]

37. Seifert, J.R.K.; Mlodzik, M. Frizzled/PCP signalling: A conserved mechanism regulating cell polarity and directed motility. *Nat. Rev. Genet.* 2007, 8, 126–138. [CrossRef]

38. Slusarski, D.C.; Corces, V.G.; Moon, R.T. Interaction of Wnt and a Frizzled homologue triggers G-protein-linked phosphatidylinositol signalling. *Nat. Cell Biol.* 1997, 390, 410–413. [CrossRef]

39. Kühl, M.; Sheldahl, L.C.; Park, M.; Miller, J.R.; Moon, R.T. The Wnt/Ca2+ pathway: A new vertebrate Wnt signaling pathway takes shape. *Trends Genet.* 2000, 16, 279–283. [CrossRef]

40. Slusarski, D.C.; Yang-Snyder, J.; Busa, W.B.; Moon, R.T. Modulation of Embryonic Intracellular Ca2+ Signaling by Wnt-5a. *Dev. Biol.* 1997, 182, 114–120. [CrossRef]

41. Sheldahl, L.C.; Park, M.; Malbon, C.C.; Moon, R.T. Protein kinase C is differentially stimulated by Wnt and Frizzled homologs in a G-protein-dependent manner. *Curr. Biol.* 1999, 9, 695–698. [CrossRef]

42. Kühl, M.; Sheldahl, L.C.; Malbon, C.C.; Moon, R.T. Ror2/Ca2+/Calmodulin-dependent Protein Kinase II Is Stimulated by Wnt and Frizzled Homologs and Promotes Ventral Cell Fates in Xenopus. *J. Biol. Chem.* 2000, 275, 12701–12711. [CrossRef] [PubMed]

43. De, A. Wnt/Ca2+ signaling pathway: A brief overview. *Acta Biochim. Biophys. Sin.* 2011, 43, 745–756. [CrossRef] [PubMed]

44. Cruciat, C.-M.; Niehrs, C. Secreted and Transmembrane Wnt Inhibitors and Activators. *Cold Spring Harb. Perspect. Biol.* 2012, 5, a015081. [CrossRef] [PubMed]

45. Semënov, M.V.; Tamaš, K.; Brott, B.K.; Kühl, M.; Sokol, S.; He, X. Head inducer Dickkopf-1 is a ligand for Wnt coreceptor LRP6. *Curr. Biol.* 2001, 11, 951–961. [CrossRef]

46. Mao, B.; Wu, W.; Davidson, G.; Marhold, J.; Li, M.; Mechler, B.M.; Delius, H.; Hoppe, D.; Stannke, P.; Walter, C.; et al. Kremen proteins are Dickkopf receptors that regulate Wnt/b-catenin signalling. *Nat. Cell Biol.* 2002, 417, 664–667. [CrossRef]

47. Mao, B.; Niehrs, C. Kremen2 modulates Dickkopf2 activity during Wnt/β-catenin signalling. *Gene* 2003, 302, 179–183. [CrossRef]

48. Hao, H.-X.; Xie, Y.; Zhang, Y.; Charlat, O.; Oster, E.; Avello, M.; Lei, H.; Mickanin, C.; Liu, D.; Ruffner, H.; et al. ZNRF3 promotes Wnt receptor turnover in an R-spondin-sensitive manner. *Nat. Cell Biol.* 2012, 485, 195–200. [CrossRef]

49. De Lau, W.; Peng, W.C.; Gros, P.; Clevers, H. The R-spondin/Lgr5/Rnf43 module: Regulator of Wnt signal strength. *Genes Dev.* 2014, 28, 305–316. [CrossRef]
50. Ohkawara, B.; Glinka, A.; Niehrs, C. Rspo3 Binds Syndecan 4 and Induces Wnt/PCP Signaling via Clathrin-Mediated Endocytosis to Promote Morphogenesis. *Dev. Cell* 2011, 20, 303–314. [CrossRef]

51. Xu, Q.; Wang, Y.; Daboub, A.; Smallwood, P.M.; Williams, J.; Woods, C.; Kelley, M.W.; Jiang, L.; Tasman, W.; Zhang, K.; et al. Vascular Development in the Retina and Inner Ear. *Cell* 2004, 116, 883–895. [CrossRef]

52. Oliva, C.A.; Montecinos-Oliva, C.; Inestrosa, N.C. Wnt Signaling in the Central Nervous System: New Insights in Health and Disease. In *Progress in Molecular Biology and Translational Science*; Elsevier: Amsterdam, The Netherlands, 2018; Volume 153, pp. 81–130.

53. Brault, V.; Moore, R.; Kutsch, S.; Ishibashi, M.; Rowitch, D.H.; McMahon, A.P.; Sommer, L.; Boussadia, O.; Kemler, R. Inactivation of the beta-catenin gene by Wnt1-Cre-mediated deletion results in dramatic brain malformation and failure of craniofacial development. *Development* 2001, 128, 1253–1264. [PubMed]

54. Thomas, K.R.; Capecchi, M.R. Targeted disruption of the murine int-1 proto-oncogene resulting in severe abnormalities in midbrain and cerebellar development. *Nat. Cell Biol.* 1990, 346, 847–850. [CrossRef] [PubMed]

55. Nordström, U.; Jessell, T.M.; Edlund, T. Progressive induction of caudal neural character by graded Wnt signaling. *Nat. Neurosci.* 2002, 5, 525–532. [CrossRef]

56. Carron, C.; Shi, D.-L. Specification of anteroposterior axis by combinatorial signaling during Xenopus development. *Wiley Interdiscip. Rev. Dev. Biol.* 2015, 5, 150–168. [CrossRef] [PubMed]

57. Glinka, A.; Wu, W.; Delius, H.; Monaghan, A.P.; Blumenstock, C.; Niehrs, C. Dickkopf-1 is a member of a new family of secreted proteins and functions in head induction. *Nat. Cell Biol.* 1998, 391, 357–362. [CrossRef] [PubMed]

58. Houart, C.; Caneparo, L.; Heisenberg, C.-P.; Barth, K.; Take-Uchi, M.; Wilson, S.W. Establishment of the Telencephalon during Gastrulation by Local Antagonism of Wnt Signaling. *Neuron* 2002, 35, 255–265. [CrossRef]

59. Zhang, X.; Cheong, S.-M.; Amado, N.G.; Reis, A.H.; Macdonald, B.T.; Zebisch, M.; Jones, E.Y.; Abreu, J.G.; He, X. Notum Is Required for Neural and Head Induction via Wnt Deacylation, Oxidation, and Inactivation. *Dev. Cell* 2015, 32, 719–730. [CrossRef] [PubMed]

60. Grove, E.A.; Tole, S.; Limon, J.; Yip, L.; Ragsdale, C.W. The hem of the embryonic cerebral cortex is defined by the expression of multiple Wnt genes and is compromised in Gli3-deficient mice. *Development* 1998, 125, 2315–2325.

61. Lee, S.M.; Tole, S.; Grove, E.; McMahon, A.P. A local Wnt-3a signal is required for development of the mammalian hippocampus. *Development* 2000, 127, 457–467.

62. Yoshinaga, Y.; Kagawa, T.; Shimizu, T.; Inoue, T.; Takada, S.; Kuratsu, J.-I.; Taga, T. Wnt3a Promotes Hippocampal Neurogenesis by Shortening Cell Cycle Duration of Neural Progenitor Cells. *Cell. Mol. Neurobiol.* 2010, 30, 1049–1058. [CrossRef]

63. Galceran, J.; Miyashita-Lin, E.M.; Devaney, E.; Rubenstein, J.L.; Grosschedl, R. Hippocampus development and generation of dentate gyrus granule cells is regulated by LEF1. *Development* 2000, 127, 469–482. [PubMed]

64. Hirabayashi, Y.; Gotoh, Y. Stage-dependent fate determination of neural precursor cells in mouse forebrain. *Neurosci. Res.* 2005, 51, 331–336. [CrossRef] [PubMed]

65. Zechnher, D.; Fujita, Y.; Hülsken, J.; Müller, T.; Walther, I.; Taketo, M.M.; Crenshaw, E.B.; Birchmeier, W.; Birchmeier, C. β-Catenin signals regulate cell growth and the balance between progenitor cell expansion and differentiation in the nervous system. *Dev. Biol.* 2003, 258, 406–418. [CrossRef]

66. Zhang, X.; Zhu, J.; Yang, G.-Y.; Wang, Q.-J.; Qian, L.; Chen, Y.-M.; Chen, F.; Tao, Y.; Hu, H.-S.; Wang, T.; et al. Dishevelled promotes axon differentiation by regulating atypical protein kinase C. *Nat. Cell Biol.* 2007, 9, 743–754. [CrossRef] [PubMed]

67. Slater, P.G.; Ramirez, V.T.; Gonzalez-Billault, C.; Varela-Nallar, L.; Inestrosa, N.C. Frizzled-5 Receptor Is Involved in Neuronal Polarity and Morphogenesis of Hippocampal Neurons. *PLoS ONE* 2013, 8, e78892. [CrossRef]

68. Ohno, S. Extrinsic Wnt signalling controls the polarity component aPKC. *Nat. Cell Biol.* 2007, 9, 738–740. [CrossRef]

69. Rosso, S.B.; Sussman, D.; Wynshaw-Boris, A.; Salinas, P.C. Wnt signaling through Dishevelled, Rac and JNK regulates dendritic development. *Nat. Neurosci.* 2005, 8, 34–42. [CrossRef]
70. Viale, B.; Song, L.; Petrenko, V.; Combremont, A.-L.W.; Contestabile, A.; Bocchi, R.; Salmon, P.; Carleton, A.; An, L.; Vutskits, L.; et al. Transient Deregulation of Canonical Wnt Signaling in Developing Pyramidal Neurons Leads to Dendritic Defects and Impaired Behavior. Cell Rep. 2019, 27, 1487–1502.e6. [CrossRef]

71. Ahmad-Annuar, A.; Ciani, L.; Simeonidis, I.; Herreros, J.; Ben Fredj, N.; Rosso, S.B.; Hall, A.; Brickley, S.; Salinas, P.C. Signaling across the synapse: A role for Wnt and Dishevelled in presynaptic assembly and neurotransmitter release. J. Cell Biol. 2006, 174, 127–139. [CrossRef]

72. Cerpa, W.; Godoy, J.A.; Alfaro, I.; Farias, G.G.; Metcalfe, M.J.; Fuenteaiba, R.; Bonansco, C.; Inestrosa, N.C. Wnt7a Modulates the Synaptic Vesicle Cycle and Somatic Synaptic Transmission in Hippocampal Neurons. J. Biol. Chem. 2007, 282, 5918–5927. [CrossRef]

73. Davis, E.K.; Zou, Y.; Ghosh, A. Wnts acting through canonical and noncanonical signaling pathways exert opposite effects on hippocampal synapse formation. Neural Dev. 2008, 3, 32. [CrossRef] [PubMed]

74. Varela-Nallar, L.; Grabowski, C.P.; Alfaro, I.E.; Alvarez, A.; Inestrosa, N.C. Role of the Wnt receptor Frizzled-1 in presynaptic differentiation and function. Neural Dev. 2009, 4, 41. [CrossRef] [PubMed]

75. Sahores, M.; Gibb, A.; Salinas, P.C. Frizzled-5, a receptor for the synaptic organizer Wnt7a, regulates activity-mediated synaptogenesis. Development 2010, 137, 2215–2225. [CrossRef]

76. Varela-Nallar, L.; Alfaro, I.E.; Serrano, F.G.; Parodi, J.; Inestrosa, N.C. Wingless-type family member 5A (Wnt-5a) stimulates synaptic differentiation and function of glutamatergic synapses. Proc. Natl. Acad. Sci. USA 2010, 107, 21164–21169. [CrossRef] [PubMed]

77. Farías, G.G.; Alfaro, I.E.; Cerpa, W.; Grabowski, C.P.; Godoy, J.A.; Bonansco, C.; Inestrosa, N.C. Wnt-5a/NF Signaling Promotes the Clustering of PSD-95 in Hippocampal Neurons. J. Biol. Chem. 2009, 284, 15857–15866. [CrossRef] [PubMed]

78. Ramírez, V.T.; Ramos-Fernández, E.; Henríquez, J.P.; Lorenzo, A.; Inestrosa, N.C. Wnt-5a/Frizzled9 Receptor Signaling through the Gαo-Gβγ Complex Regulates Dendritic Spine Formation. J. Biol. Chem. 2016, 291, 19092–19107. [CrossRef] [PubMed]

79. Ciani, L.; Boyle, K.A.; Dickins, E.; Sahores, M.; Anane, D.; Lopes, D.M.; Gibb, A.J.; Salinas, P.C. Wnt7a signaling promotes dendritic spine growth and synaptic strength through Ca2+/Calmodulin-dependent protein kinase II. Proc. Natl. Acad. Sci. USA 2011, 108, 10732–10737. [CrossRef] [PubMed]

80. Oliva, C.A.; Inestrosa, N.C. A novel function for Wnt signaling modulating neuronal firing activity and the temporal structure of spontaneous oscillation in the entorhinal–hippocampal circuit. Exp. Neurol. 2015, 269, 43–55. [CrossRef]

81. Cerpa, W.; Gambrill, A.; Inestrosa, N.C.; Barria, A. Regulation of NMDA-Receptor Synaptic Transmission by Wnt Signaling. J. Neurosci. 2011, 31, 9466–9471. [CrossRef]

82. Gogolla, N.; Galimberti, I.; Deguchi, Y.; Caroni, P. Wnt Signaling Mediates Experience-Related Regulation of Synapse Numbers and Mossy Fiber Connectivities in the Adult Hippocampus. Neuron 2009, 62, 510–525. [CrossRef]

83. McLeod, F.; Salinas, P.C. Wnt proteins as modulators of synaptic plasticity. Curr. Opin. Neurobiol. 2018, 53, 90–95. [CrossRef] [PubMed]

84. McLeod, F.; Bossio, A.; Marzo, A.; Ciani, L.; Sibilla, S.; Hannan, S.; Wilson, G.A.; Palomer, E.; Smart, T.G.; Gibb, A.; et al. Wnt Signaling Mediates LTP-Dependent Spine Plasticity and AMPAR Localization through Frizzled-7 Receptors. Cell Rep. 2018, 23, 1060–1071. [CrossRef] [PubMed]

85. Ciani, L.; Marzo, A.; Boyle, K.; Stamatakou, E.; Lopes, D.M.; Anane, D.; McLeod, F.; Rosso, S.B.; Gibb, A.; Salinas, P.C. Wnt signalling tunes neurotransmitter release by directly targeting Synaptotagmin-1. Nat. Commun. 2015, 6, 8302. [CrossRef] [PubMed]

86. Chen, J.; Park, C.S.; Tang, S.-J. Activity-dependent Synaptic Wnt Release Regulates Hippocampal Long Term Potentiation. J. Biol. Chem. 2006, 281, 11910–11916. [CrossRef] [PubMed]

87. Vargas, J.Y.; Fuenzalida, M.; Inestrosa, N.C. In vivo Activation of Wnt Signaling Pathway Enhances Cognitive Function of Adult Mice and Reverses Cognitive Deficits in an Alzheimer’s Disease Model. J. Neurosci. 2014, 34, 2191–2202. [CrossRef] [PubMed]

88. Tabatadze, N.; Tomas, C.; McGonigal, R.; Lin, B.; Schook, A.; Routtenberg, A. Wnt transmembrane signaling and long-term spatial memory. Hippocampus 2011, 22, 1228–1241. [CrossRef] [PubMed]

89. Fortress, A.M.; Schram, S.L.; Tuscher, J.J.; Frick, K.M. Canonical Wnt Signaling is Necessary for Object Recognition Memory Consolidation. J. Neurosci. 2013, 33, 12619–12626. [CrossRef] [PubMed]
90. Gonçalves, J.T.; Schafer, S.T.; Gage, F.H. Adult Neurogenesis in the Hippocampus: From Stem Cells to Behavior. *Cell* 2016, 167, 897–914. [CrossRef] [PubMed]

91. Kumar, A.; Pareek, V.; Faiq, M.A.; Ghosh, S.K.; Kumari, C. ADULT NEUROGENESIS IN HUMANS: A Review of Basic Concepts, History, Current Research, and Clinical Implications. *Curr. Res. Clin. Implic. Innov. Clin. Neurosci.* 2019, 16, 30–37.

92. Lie, D.C.; Colamarino, S.A.; Song, H.-J.; Désiré, L.; Mira, H.; Consiglio, A.; Lein, E.S.; Jessberger, S.; Lansford, H.; Dearie, A.R.; et al. Wnt signalling regulates adult hippocampal neurogenesis. *Nat. Cell Biol.* 2005, 437, 1370–1375. [CrossRef] [PubMed]

93. Jang, M.-H.; Bonaguidi, M.A.; Kitabatake, Y.; Sun, J.; Song, J.; Kang, E.; Jun, H.; Zhong, C.; Su, Y.; Guo, J.U.; et al. Secreted Frizzled-Related Protein 3 Regulates Activity-Dependent Adult Hippocampal Neurogenesis. *Cell Stem Cell* 2013, 12, 215–223. [CrossRef] [PubMed]

94. Jessberger, S.; Clark, R.E.; Broadbent, N.J.; Clemenson, J.G.D.; Consiglio, A.; Lie, D.C.; Squire, L.R.; Gage, F.H. Dentate gyrus-specific knockdown of adult neurogenesis impairs spatial and object recognition memory in adult rats. *Learn. Mem.* 2009, 16, 147–154. [CrossRef] [PubMed]

95. Schwab, M.H.; Bartholomaeus, A.; Heinrich, B.; Feldmeyer, D.; Druffel-Augustin, S.; Goebbels, S.; Naya, F.J.; Zhao, S.; Frotscher, M.; Tsai, M.-J.; et al. Neuronal Basic Helix-Loop-Helix Proteins (NEX and BETA2/Neuro D) Regulate Terminal Granule Cell Differentiation in the Hippocampus. *J. Neurosci.* 2000, 20, 3714–3724. [CrossRef] [PubMed]

96. Kuwabara, T.; Hsieh, J.; Muotri, A.R.; Yeo, G.; Warashina, M.; Lie, D.C.; Moore, L.; Nakashima, K.; Asashima, M.; Gage, F.H. Wnt-mediated activation of NeuroD1 and retro-elements during adult neurogenesis. *Nat. Neurosci.* 2009, 12, 1097–1105. [CrossRef] [PubMed]

97. Qu, Q.; Sun, G.; Murai, K.; Ye, P.; Li, W.; Asuelime, G.; Cheung, Y.-T.; Shi, Y. Wnt7a Regulates Multiple Steps in rat hippocampus. *Aging Cell* 2017, 16, 1195–1199. [CrossRef] [PubMed]

98. García-Velázquez, L.; Arias, C. The emerging role of Wnt signaling dysregulation in the understanding and modification of age-associated diseases. *Aging Res. Rev.* 2017, 37, 135–145. [CrossRef]

99. Tobin, M.K.; Musaraca, K.; Disouky, A.; Shetti, A.; Bheri, A.; Honer, W.G.; Kim, N.; Dawe, R.J.; Bennett, D.A.; Arfanakis, K.; et al. Human Hippocampal Neurogenesis Persists in Aged Adults and Alzheimer’s Disease Patients. *Cell Stem Cell* 2019, 24, 974–982.e3. [CrossRef]

100. Mathews, K.J.; Allen, K.M.; Boerrigter, D.; Ball, H.; Weickert, C.S.; Double, K.L. Evidence for reduced neurogenesis in the aging human hippocampus despite stable stem cell markers. *Aging Cell* 2017, 16, 1195–1199. [CrossRef] [PubMed]

101. Petralia, R.S.; Mattson, M.P.; Yao, P.J. Communication breakdown: The impact of ageing on synapse structure. *Aging Res. Rev.* 2014, 14, 31–42. [CrossRef]

102. Morrison, J.H.; Baxter, M.G. The ageing cortical synapse: Hallmarks and implications for cognitive decline. *Nat. Rev. Neurosci.* 2012, 13, 240–250. [CrossRef] [PubMed]

103. Hofmann, J.W.; McBryan, T.; Adams, P.D.; Sedivy, J.M. The effects of aging on the expression of Wnt pathway genes in mouse tissues. *AGE* 2014, 36, 1033–1040. [CrossRef] [PubMed]

104. Okamoto, M.; Inoue, K.; Iwamura, H.; Terashima, K.; Soya, H.; Asashima, M.; Kuwabara, T. Reduction in paracrine Wnt3 factors during aging causes impaired adult neurogenesis. *FASEB J.* 2011, 25, 3570–3582. [CrossRef] [PubMed]

105. Seib, D.R.; Corsini, N.S.; Ellwanger, K.; Plaa, C.; Mateos, A.; Pitzer, C.; Niehrs, C.; Celikel, T.; Martin-Villalba, A. Loss of Dickkopf-1 Restores Neurogenesis in Old Age and Counteracts Cognitive Decline. *Cell Stem Cell* 2013, 12, 204–214. [CrossRef]

106. Orellana, A.M.M.; Vasconcelos, A.R.; Leite, J.A.; Lima, L.D.S.; Andreotti, D.Z.; Munhoz, C.D.; Kawamoto, E.M.; Scavone, C. Age-related neuroinflammation and changes in AKT-GSK-3β and WNT/β-CATENIN signaling in rat hippocampus. *Aging* 2015, 7, 1094–1108. [CrossRef]

107. Folke, J.; Pakkenberg, B.; Brudek, T. Impaired Wnt Signaling in the Prefrontal Cortex of Alzheimer’s Disease. *Mol. Neurobiol.* 2018, 56, 873–891. [CrossRef]
109. Brann, D.W.; Dhandapani, K.; Wakade, C.; Mahesh, V.B.; Khan, M.M. Neurotrophic and neuroprotective actions of estrogen: Basic mechanisms and clinical implications. *Steroids* 2007, 72, 381–405. [CrossRef]

110. Scott, E.L.; Zhang, Q.-G.; Han, N.; Desai, B.N.; Brann, D.W. Long-term estrogen deprivation leads to elevation of Dickkopf-1 and dysregulation of Wnt/β-Catenin signaling in hippocampal CA1 neurons. *Steroids* 2013, 78, 624–632. [CrossRef]

111. Marzo, A.; Galli, S.; Lopes, D.; McLeod, F.; Podpolny, M.; Segovia-Roldan, M.; Ciani, L.; Purro, S.; Cacucci, F.; Gibb, A.; et al. Reversal of Synapse Degeneration by Restoring Wnt Signaling in the Adult Hippocampus. *Curr. Biol.* 2016, 26, 2551–2561. [CrossRef]

112. Anderton, B.H.; Dayanandan, R.; Killlick, R.; Lovestone, S. Does dysregulation of the Notch and wingless/Wnt pathways underlie the pathogenesis of Alzheimer’s disease? *Mol. Med. Today* 2000, 6, 54–59. [CrossRef]

113. Van Der Flier, W.M. Epidemiology and risk factors of dementia. *J. Neurosci.* 2006, 228–232. [CrossRef] [PubMed]

114. Mateo, I.; Infante, J.; Llorca, J.; Rodríguez, E.; Berciano, J.; Combarros, O. Association between Glycogen Synthase Kinase-3β Genetic Polymorphism and Late-Onset Alzheimer’s Disease. *Dement. Geriatr. Cogn. Disord.* 2006, 21, 228–232. [CrossRef] [PubMed]

115. De Ferrari, G.V.; Papassotiropoulos, A.; Biechele, T.; De-Vrieze, F.W.; Avila, M.E.; Major, M.B.; Myers, A.; Sáez, K.; Henriquez, J.P.; Zhao, A.; et al. Common genetic variation within the Low-Density Lipoprotein Receptor-Related Protein 6 and late-onset Alzheimer’s disease. *Proc. Natl. Acad. Sci. USA* 2007, 104, 9434–9439. [CrossRef] [PubMed]

116. Alarcón, M.A.; Medina, M.A.; Hu, Q.; Avila, M.E.; Bustos, B.I.; Pérez-Palma, E.; Peralta, A.; Salazar, P.; Ugarte, G.D.; Reyes, A.E.; et al. A novel functional low-density lipoprotein receptor-related protein 6 gene alternative splice variant is associated with Alzheimer’s disease. *Neurobiol. Aging* 2013, 34, 1709.e9–1709.e18. [CrossRef] [PubMed]

117. Liu, C.-C.; Tsai, C.-W.; Deak, F.; Rogers, J.; Penuliar, M.; Sung, Y.M.; Maher, J.N.; Fu, Y.; Li, X.; Xu, H.; et al. Deficiency in LRP6-Mediated Wnt Signaling Contributes to Synaptic Abnormalities and Amyloid Pathology in Alzheimer’s Disease. *Neuron* 2014, 84, 63–77. [CrossRef]

118. Buechler, J.; Salinas, P.C. Deficient Wnt Signaling and Synaptic Vulnerability in Alzheimer’s Disease: Emerging Roles for the LRP6 Receptor. *Front. Synaptic Neurosci.* 2018, 10, 38. [CrossRef]

119. Caricasole, A.; Copani, A.; Caraci, F.; Aronica, E.; Rozemuller, A.J.; Caruso, A.; Storto, M.; Gaviraghi, G.; Terstappen, G.C.; Nicoletti, F. Induction of Dickkopf-1, a Negative Modulator of the Wnt Pathway, Is Associated with Neuronal Degeneration in Alzheimer’s Brain. *J. Neurosci.* 2004, 24, 6021–6027. [CrossRef]

120. Rosi, M.C.; Luccarini, I.; Grossi, C.; Fiorentini, A.; Spillantini, M.G.; Prisco, A.; Scali, C.; Gianfriddo, M.; Caricasole, A.; Terstappen, G.C.; et al. Increased Dickkopf-1 expression in transgenic mouse models of neurodegenerative disease. *J. Neurochem.* 2010, 112, 1539–1551. [CrossRef]

121. Purro, S.A.; Dickins, E.M.; Salinas, P.C. The Secreted Wnt Antagonist Dickkopf-1 Is Required for Amyloid-Mediated Synaptic Loss. *J. Neurosci.* 2012, 32, 3492–3498. [CrossRef]

122. Friedman, B.A.; Srinivasan, K.; Ayalon, G.; Meilandt, W.J.; Lin, H.; Huntley, M.A.; Cao, Y.; Lee, S.-H.; Haddick, P.C.; Ngu, H.; et al. Diverse Brain Myeloid Expression Profiles Reveal Distinct Microglial Activation States and Aspects of Alzheimer’s Disease Not Evident in Mouse Models. *Cell Rep.* 2018, 22, 832–847. [CrossRef]

123. Lambert, J.-C.; The European Alzheimer’s Disease Initiative Investigators; Heath, S.; Even, G.; Campion, D.; Sleebers, K.; Hiltunen, M.O.; Combarros, O.; Zelenika, D.; Bullido, M.J.; et al. Genome-wide association study identifies variants at CLU and CR1 associated with Alzheimer’s disease. *Nat. Genet.* 2009, 41, 1094–1099. [CrossRef] [PubMed]

124. Harold, D.; Abraham, R.; Hollingworth, P.; Sims, R.; Gerrish, A.; Hamshere, M.L.; Patel, J.; Moskvina, V.; Dowzell, K.; Williams, A.; et al. Genome-wide association study identifies variants at CLU and PICALM associated with Alzheimer’s disease. *Nat. Genet.* 2009, 41, 1088–1093. [CrossRef] [PubMed]

125. Liu, C.-C.; Kanekiyo, T.; Xu, H.; Bu, G. Apolipoprotein E and Alzheimer disease: Risk, mechanisms and therapy. *Nat. Rev. Neurosci.* 2013, 9, 106–118. [CrossRef]

126. Bu, G. Apolipoprotein E and its receptors in Alzheimer’s disease: Pathways, pathogenesis and therapy. *Nat. Rev. Neurosci.* 2009, 10, 333–344. [CrossRef] [PubMed]
127. Farrer, L.A.; Cupples, L.A.; Haines, J.L.; Hyman, B.T.; Kukull, W.A.; Mayeux, R.; Myers, R.H.; Pericak-Vance, M.A.; Risch, N.; Van Duijn, C.M. Effects of Age, Sex, and Ethnicity on the Association Between Apolipoprotein E Genotype and Alzheimer Disease. *JAMA* 1997, 278, 1349–1356. [CrossRef] [PubMed]

128. Schmechel, D.E.; Saunders, A.M.; Strittmatter, W.J.; Crain, B.J.; Hulette, C.M.; Joo, S.H.; Pericak-Vance, M.A.; Goldgaber, D.; Roses, A.D. Increased amyloid beta-peptide deposition in cerebral cortex as a consequence of apolipoprotein E genotype in late-onset Alzheimer disease. *Proc. Natl. Acad. Sci. USA* 1993, 90, 9649–9653. [CrossRef] [PubMed]

129. Tachibana, M.; Holm, M.-L.; Liu, C.-C.; Shinohara, M.; Aikawa, T.; Oue, H.; Yamazaki, Y.; Martens, Y.A.; Murray, M.E.; Sullivan, P.M.; et al. APOE4-mediated amyloid-β pathology depends on its neuronal receptor LRP1. *J. Clin. Investig.* 2019, 129, 1272–1277. [CrossRef]

130. Zilberberg, A.; Yaniv, A.; Gazit, A. The Low Density Lipoprotein Receptor-1, LRP1, Interacts with the Human Frizzled-1 (HFrz1) and Down-regulates the Canonical Wnt Signaling Pathway. *J. Biol. Chem.* 2004, 279, 17535–17542. [CrossRef]

131. Jonsson, T.; Stefansson, H.; Steinberg, S.; Jonsdottir, I.; Jonsson, P.V.; Snaedal, J.; Bjornsson, S.; Caruso, A.; Motolese, M.; Iacovelli, L.; Caraci, F.; Copani, A.; Nicoletti, F.; Terstappen, G.C.; Gaviraghi, G.; Caricasole, A. Inhibition of the canonical Wnt signaling pathway by apolipoprotein E4 in PC12 cells. *J. Neurochem.* 2006, 98, 364–371. [CrossRef]

132. Yue, W.; Sun, Q.; Dacic, S.; Landreneau, R.J.; Siegfried, J.M.; Yu, J.; Zhang, L. Downregulation of Dkk3. *BMC Cell Biol.* 2018, 19, 8, 1–13. [CrossRef]

133. Xu, J.; Patassini, S.; Rustogi, N.; Riba-Garcia, I.; Hale, B.D.; Phillips, A.M.; Waldvogel, H.; Haines, R.; Bradbury, P.; Stevens, A.; et al. Regional protein expression in human Alzheimer’s brain correlates with disease severity. *Commun. Biol.* 2019, 2, 1–15. [CrossRef]

134. Elliott, C.; Rojo, A.I.; Ribe, E.; Broadstock, M.; Xia, W.; Morin, P.; Semenov, M.; Baillie, G.; Cuadrado, A.; Caricasole, A.; Romeo, A.; Vital, A.; Dumanchin, C.; et al. TREM2 Promotes Microglial Survival by Activating Wnt/β-Catenin Pathway. *J. Neurosci.* 2017, 37, 1772–1784. [CrossRef] [PubMed]

135. Bai, B.; Wang, X.; Li, Y.; Chen, P.-C.; Yu, K.; Dey, K.K.; Yarbro, J.M.; Han, X.; Lutz, B.M.; Rao, S.; et al. Deep Multilayer Brain Proteomics Identifies Molecular Networks in Alzheimer’s Disease’s Progression. *Neuron* 2020, 105, 975–991.e7. [CrossRef] [PubMed]

136. Hooi, B.; Tanzi, R.E. The Genetic Basis of Alzheimer’s Disease: Findings From Genome-Wide Studies. In *Genomics, Circuits, and Pathways in Clinical Neuropsychiatry*; Elsevier Inc.: Amsterdam, The Netherlands, 2016; pp. 547–571.

137. Sleegers, K.; Brouwers, N.; Gijselink, I.; Theuns, J.; Goossens, D.; Wauters, J.; Del-Favero, J.; Cruts, M.; Van Duijn, C.M.; Van Broeckhoven, C. APP duplication is sufficient to cause early onset Alzheimer’s dementia with cerebral amyloid angiopathy. *Brain* 2006, 129, 2977–2983. [CrossRef]

138. Rovelet-Lecrux, A.; Hannequin, D.; Raux, G.; Le Meur, N.; Laquerrière, A.; Vital, A.; Dumanchin, C.; Feuillette, S.; Brice, A.; Vercelletto, M.; et al. APP locus duplication causes autosomal dominant early-onset Alzheimer disease with cerebral amyloid angiopathy. *Nat. Genet.* 2006, 38, 24–26. [CrossRef] [PubMed]
Brain Sci. 2020, 10, 902

145. De Strooper, B.; Saftig, P.; Craessaerts, K.; Vanderstichele, H.; Guhde, G.; Annaert, W.; Von Figura, K.; Van Leuven, F. Deficiency of presenilin-1 inhibits the normal cleavage of amyloid precursor protein. Nat. Cell Biol. 1998, 391, 387–390. [CrossRef] [PubMed]

146. Zhang, N.; Parr, C.J.; Birch, A.M.; Goldfinger, M.H.; Sastre, M. The amyloid precursor protein binds to β-catenin and modulates its cellular distribution. Neurosci. Lett. 2018, 685, 190–195. [CrossRef] [PubMed]

147. Zhang, Z.; Hartmann, H.; Do, V.M.; Abramowski, D.; Sturchler-Pierrat, C.; Staufenbiel, M.; Sommer, B.; Van De Wetering, M.; Clevers, H.; Saftig, P.; et al. Destabilization of β-catenin by mutations in presenilin-1 potentiates neuronal apoptosis. Nat. Cell Biol. 1998, 395, 698–702. [CrossRef]

148. Nishimura, M.; Yu, G.; Levesque, G.; Zhang, D.; Ruel, L.; Chen, F.; Milman, P.; Holmes, E.; Liang, Y.; Kawarai, T.; et al. Presenilin mutations associated with Alzheimer disease cause defective intracellular trafficking of β-catenin, a component of the presenilin protein complex. Nat. Med. 1999, 5, 164–169. [CrossRef]

149. Kang, D.E.; Soriano, S.; Frosch, M.P.; Collins, T.; Naruse, S.; Sisodia, S.S.; Leibowitz, G.; Levine, F.; Van Leuven, F. Deficiency of presenilin-1 inhibits the normal cleavage of amyloid precursor protein. Acta Neuropathol. 2000, 90, 27 of 33

150. Teo, J.-L.; Ma, H.; Nguyen, C.; Lam, C.; Kahn, M. Specific inhibition of CBP/β-catenin interaction rescues defects in neuronal differentiation caused by a presenilin-1 mutation. Proc. Natl. Acad. Sci. USA 2005, 102, 12171–12176. [CrossRef]

151. Boonen, R.A.; Van Tijn, P.; Zivkovic, D. Wnt signaling in Alzheimer’s disease: Up or down, that is the question. Ageing Res. Rev. 2009, 8, 71–82. [CrossRef]

152. Tapia-Rojas, C.; Burgos, P.V.; Inestrosa, N.C. Inhibition of Wnt signaling induces amyloidogenic processing of amyloid precursor protein and the production and aggregation of Amyloid-β (Aβ)42 peptides. J. Neurochem. 2016, 139, 1175–1191. [CrossRef]

153. Parr, C.; Mirzaei, N.; Christian, M.; Sastre, M. Activation of the Wnt/β-catenin pathway represses the transcription of the β-amyloid precursor protein cleaving enzyme (BACE1) via binding of T-cell factor-4 to BACE1 promoter. FASEB J. 2014, 29, 623–635. [CrossRef]

154. Vargas, J.Y.; Ahumada, J.; Arráezola, M.S.; Fuenzalida, M.; Inestrosa, N.C. WASP-1, a canonical Wnt signaling potentiator, rescues hippocampal synaptic impairments induced by Aβ oligomers. Exp. Neurol. 2015, 264, 14–25. [CrossRef] [PubMed]

155. Tapia-Rojas, C.; Inestrosa, N.C. Wnt signaling loss accelerates the appearance of neuropathological hallmarks of Alzheimer’s disease in J20-APP transgenic and wild-type mice. J. Neurochem. 2018, 144, 443–465. [CrossRef] [PubMed]

156. De Ferrari, G.V.; Chacon, M.A.; Barria, M.I.; Garrido, J.L.; Godoy, J.A.; Olivares, G.H.; Reyes, A.E.; Alvarez, A.R.; Bronfman, M.; Inestrosa, N.C. Activation of Wnt signaling rescues neurodegeneration and behavioral impairments induced by β-amyloid fibrils. Mol. Psychiatry 2003, 8, 195–208. [CrossRef] [PubMed]

157. Alvarez, A.R.; Godoy, J.A.; Mullendorff, K.; Olivares, G.H.; Bronfman, M.; Inestrosa, N.C. Wnt-3a overcomes β-amyloid toxicity in rat hippocampal neurons. Exp. Cell Res. 2004, 297, 186–196. [CrossRef]

158. Quintanilla, R.A.; Muñoz, F.J.; Metcalfe, M.J.; Hitzfeld, M.; Olivares, G.; Godoy, J.A.; Inestrosa, N.C. Trolax and 17-Estradiol Protect against Amyloid-β Peptide Neurotoxicity by a Mechanism That Involves Modulation of the Wnt Signaling Pathway. J. Biol. Chem. 2005, 280, 11615–11625. [CrossRef]

159. Magdesian, M.H.; Carvalho, M.M.V.F.; Mendes, F.A.; Saraiva, L.M.; Juliano, M.A.; Juliano, L.; Garcia-Abreu, J.; Ferreira, S.T. Amyloid-β Binds to the Extracellular Cysteine-rich Domain of Frizzled and Inhibits Wnt/β-Catenin Signaling. J. Biol. Chem. 2008, 283, 9359–9368. [CrossRef] [PubMed]

160. Barbier, P.; Zejneli, O.; Martinho, M.; Lasorsa, A.; Belle, V.; Smet-Noccia, C.; Tsvetkov, P.O.; Devred, F.; Landrieu, I. Role of Tau as a Microtubule-Associated Protein: Structural and Functional Aspects. Front. Aging Neurosci. 2019, 11, 204. [CrossRef]

161. Iqbal, K.; Liu, F.; Gong, C.-X.; Grundke-Iqbal, I. Tau in Alzheimer Disease and Related Tauopathies. Curr. Alzheimer Res. 2010, 7, 656–664. [CrossRef]

162. Guo, T.; Noble, W.; Hanger, D.P. Roles of tau protein in health and disease. Acta Neuropathol. 2017, 133, 665–704. [CrossRef]

163. Hutton, M.; Lendon, C.L.; Rizzu, P.; Baker, M.; Froelich, S.; Houlden, H.; Pickering-Brown, S.; Chakraverty, S.; Isaacs, A.; Grover, A.; et al. Association of missense and 5′-splice-site mutations in tau with the inherited dementia FTDP-17. Nat. Cell Biol. 1998, 393, 702–705. [CrossRef]
164. Small, S.A.; Duff, K. Linking Aβ and Tau in Late-Onset Alzheimer’s Disease: A Dual Pathway Hypothesis. *Neuron* 2008, 60, 534–542. [CrossRef] [PubMed]

165. Ittner, L.M.; Ke, Y.D.; Delerue, F.; Bi, M.; Gladbach, A.; Van Eersel, J.; Wölfing, H.; Chieng, B.C.; Christie, M.J.; Napier, L.A.; et al. Dendritic Function of Tau Mediates Amyloid-β Toxicity in Alzheimer’s Disease Mouse Models. *Cell 2010*, 142, 387–397. [CrossRef] [PubMed]

166. Roberson, E.D.; Halabisky, B.; Yoo, J.W.; Yao, J.; Chin, J.; Yan, F.; Wu, T.; Hamto, P.; Devidze, N.; Yu, G.-Q.; et al. Amyloid-β/PrP-Induced Synaptic, Network, and Cognitive Impairments Depend on Tau Levels in Multiple Mouse Models of Alzheimer’s Disease. *J. Neurosci.* 2011, 31, 700–711. [CrossRef] [PubMed]

167. Roberson, E.D.; Scearce-Levie, K.; Palop, J.J.; Yan, F.; Cheng, I.H.; Wu, T.; Gerstein, H.; Yu, G.-Q.; Mucke, L. Reducing Endogenous Tau Ameliorates Amyloid-β-Induced Deficits in an Alzheimer’s Disease Mouse Model. *Science 2007*, 316, 750–754. [CrossRef] [PubMed]

168. Pickett, E.K.; Herrmann, A.G.; McQueen, J.; Abt, K.; Dando, O.; Tulloch, J.; Jain, P.; Dunnett, S.; Sohlra, S.; Fjeldstad, M.P.; et al. Amyloid Beta and Tau Cooperate to Cause Reversible Behavioral and Transcriptional Deficits in a Model of Alzheimer’s Disease. *Cell Rep.* 2019, 29, 3592–3604.e5. [CrossRef] [PubMed]

169. Ballatore, C.; Lee, V.M.-Y.; Trojanowski, J.Q. Tau-mediated neurodegeneration in Alzheimer’s disease and related disorders. *Nat. Rev. Neurosci.* 2007, 8, 663–672. [CrossRef]

170. Busche, M.A.; Hyman, B.T. Synergy between amyloid-β and tau in Alzheimer’s disease. *Nat. Neurosci.* 2020, 23, 1183–1193. [CrossRef]

171. Weigand, A.J.; Bangen, K.J.; Thomas, K.R.; Delano-Wood, L.; Gilbert, P.E.; Brickman, A.M.; Bondi, M.W.; Pickett, E.K.; Herrmann, A.G.; McQueen, J.; Abt, K.; Dando, O.; Tulloch, J.; Jain, P.; Dunnett, S.; Sohlra, S.; Fjeldstad, M.P.; et al. Amyloid Beta and Tau Cooperate to Cause Reversible Behavioral and Transcriptional Deficits in a Model of Alzheimer’s Disease. *Cell Rep.* 2019, 29, 3592–3604.e5. [CrossRef] [PubMed]

172. Van Der Kant, R.; Goldstein, L.S.B.; Ossenkoppele, R. Amyloid-β-independent regulators of tau pathology in Alzheimer disease. *Nat. Rev. Neurosci.* 2020, 21, 21–35. [CrossRef]

173. Cavallini, A.; Brewerton, S.; Bell, A.; Sargent, S.; Glover, S.; Hardy, C.; Moore, R.; Galley, J.; Ramachandran, D.; Poidinger, M.; et al. An Unbiased Approach to Identifying Tau Kinases That Phosphorylate Tau at Sites Associated with Alzheimer Disease. *J. Biol. Chem.* 2013, 288, 23331–23347. [CrossRef]

174. Asuni, A.A.; Hooper, C.; Reynolds, C.H.; Lovestone, S.; Anderton, B.H.; Killick, R. GSK3β exhibits β-catenin and tau directed kinase activities that are modulated by Wnt. *Eur. J. Neurosci.* 2020, 24, 3387–3392. [CrossRef] [PubMed]

175. Salcedo-Tello, P.; Hernández-Ortega, K.; Arias, C. Susceptibility to GSK3β-Induced Tau Phosphorylation Differs Between the Young and Aged Hippocampus after Wnt Signaling Inhibition. *J. Alzheimer’s Dis.* 2014, 39, 775–785. [CrossRef] [PubMed]

176. Scali, C.; Caraci, F.; Gianfriddo, M.; Diodato, E.; Roncarati, R.; Pollio, G.; Gaviraghi, G.; Copani, A.; Nicoletti, F.; Terstappen, G.C.; et al. Inhibition of Wnt signaling, modulation of Tau phosphorylation and induction of neuronal cell death by DKK1. *Neurobiol. Dis.* 2006, 24, 254–265. [CrossRef] [PubMed]

177. Garrido, J.L.; Godoy, J.A.; Alvarez, A.; Bronfman, M.; Inestrosa, N.C. Protein kinase C inhibits amyloid-β peptide neurotoxicity by acting on members of the Wnt pathway. *FASEB J.* 2002, 16, 1982–1984. [CrossRef]

178. Hooper, C.; Killick, R.; Lovestone, S. The GSK3β hypothesis of Alzheimer’s disease. *J. Neurochem.* 2008, 104, 1433–1439. [CrossRef] [PubMed]

179. Llorens-Martín, M.; Jurado, J.; Hernández, F.; Ávila, J. GSK-3β, a pivotal kinase in Alzheimer disease. *Front. Mol. Neurosci.* 2014, 7, 46. [CrossRef] [PubMed]

180. Takashima, A.; Murayama, M.; Murayama, O.; Kohno, T.; Honda, T.; Yasutake, K.; Nihonmatsu, N.; Mercken, M.; Yamaguchi, H.; Sugihara, S.; et al. Presenilin 1 associates with glycogen synthase kinase-3β and its substrate tau. *Proc. Natl. Acad. Sci. USA* 1998, 95, 9637–9641. [CrossRef]

181. Cohen, T.J.; Friedmann, D.; Hwang, A.W.; Marmorstein, R.; Lee, V.M.Y. The microtubule-associated tau protein has intrinsic acetyltransferase activity. *Nat. Struct. Mol. Biol.* 2013, 20, 756–762. [CrossRef]

182. Liu, E.; Zhou, Q.; Xie, A.; Li, X.; Li, M.; Ye, J.; Li, S.; Ke, D.; Wang, Q.; Xu, Z.; et al. Tau acetylates and stabilizes β-catenin thereby promoting cell survival. *EMBO Rep.* 2020, 21, e48328. [CrossRef]

183. Sperling, R.A.; Aisen, P.S.; Beckett, L.A.; Bennett, D.A.; Craft, S.; Fagan, A.M.; Iwatsubo, T.; Jack, C.R., Jr.; Kaye, J.; Montine, T.J.; et al. Toward defining the preclinical stages of Alzheimer’s disease: Recommendations from the National Institute on Aging-Alzheimer’s Association workgroups on diagnostic guidelines for Alzheimer’s disease. *Alzheimer’s Dement.* 2011, 7, 280–292. [CrossRef]
Brain Sci. 2020, 10, 902

184. Rodriguez, J.L.; Karikari, T.K.; Suárez-Calvet, M.; Troakes, C.; King, A.; Emersic, A.; Aarsland, D.; Hye, A.; Zetterberg, H.; Blennow, K.; et al. Plasma p-tau181 accurately predicts Alzheimer’s disease pathology at least 8 years prior to post-mortem and improves the clinical characterisation of cognitive decline. Acta Neuropathol. 2020, 140, 267–278. [CrossRef] [PubMed]

185. Shankar, G.M.; Bloodgood, B.L.; Townsend, M.; Walsh, D.M.; Selkoe, D.; Sabatini, B.L. Natural Oligomers of the Alzheimer Amyloid-Protein Induce Reversible Synapse Loss by Modulating an NMDA-Type Glutamate Receptor-Dependent Signaling Pathway. J. Neurosci. 2007, 27, 2866–2875. [CrossRef] [PubMed]

186. Lacor, P.N.; Buniel, M.C.; Furlow, P.W.; Clemente, A.S.; Velasco, P.T.; Wood, M.; Viola, K.L.; Klein, W.L. Synaptic transmission is impaired prior to plaque formation in amyloid precursor protein-overexpressing mice without altering behaviorally-correlated sharp wave–ripple complexes. Neuroscience 2009, 162, 1081–1090. [CrossRef]

187. Selkoe, D. Alzheimer’s Disease Is a Synaptic Failure. Science 2002, 298, 789–791. [CrossRef] [PubMed]

188. Hsia, A.Y.; Masliah, E.; McConlogue, L.; Yu, G.-Q.; Tatsuno, G.; Hu, K.; Malenka, R.C.; Kremen1. et al. Amyloid-β-Oligomer-Induced Synapse Loss in Neuronal Cell Culture Model of Alzheimer’s Disease by Silencing Apolipoprotein E4 Carriers. Acta Neuropathol. 2018, 135, 87. [CrossRef] [PubMed]

189. Hermann, D.; Both, M.; Ebert, U.; Gross, G.; Schoemaker, H.; Draguhn, A.; Wicke, K.; Nimmrich, V. Synaptic transmission is impaired prior to plaque formation in amyloid precursor protein–overexpressing mice—without altering behaviorally-correlated sharp wave–ripple complexes. Neuroscience 2009, 162, 1081–1090. [CrossRef]

190. Cerpa, W.; Farias, G.G.; Godoy, J.A.; Fuenzalida, M.; Bonansco, P.; Inestrosa, N.C. Wnt-5a occludes Aβ oligomer-induced depression of glutamatergic transmission in hippocampal neurons. Mol. Neurodegener. 2010, 5, 3. [CrossRef]

191. DeKosky, S.T.; Scheff, S.W. Synapse loss in frontal cortex biopsies in Alzheimer’s disease: Correlation with cognitive severity. Ann. Neurol. 1990, 27, 457–464. [CrossRef]

192. Scheff, S.W.; Price, D.A.; Schmitt, F.A.; Muñson, E.J. Hippocampal synaptic loss in early Alzheimer’s disease and mild cognitive impairment. Neuropsychol. Aging 2006, 27, 1372–1384. [CrossRef]

193. Terry, R.D.; Masliah, E.; Salmon, D.P.; Butters, N.; Be, R.D.; Hill, R.; Hansen, L.A.; Katzman, R. Physical basis of cognitive alterations in Alzheimer’s disease: Synapse loss is the major correlate of cognitive impairment. Ann. Neurol. 1991, 30, 572–580. [CrossRef]

194. Forner, S.; Baglietto-Vargas, D.; Martini, A.C.; Trujillo-Estrada, L.; LaFerla, F.M. Deficient Wnt signalling triggers synaptotoxicity is Wnt-PCP dependent and blocked by fasudil. Mol. Neurodegener. 2020, 18, 3. [CrossRef] [PubMed]

195. Killick, R.; Ribe, E.M.; Alshawi, R.; Malik, B.; Hoppe, D.; Fernandes, C.P.D.; Dobson, R.; Nolan, P.M.; Lourdusamy, A.; Furney, S.J.; et al. Clusterin regulates β-amyloid toxicity via Dickkopf-1-driven induction of the wnt–PCP–JNK pathway. Mol. Psychiatry 2014, 19, 88–98. [CrossRef] [PubMed]

196. Jackson, R.J.; Rose, J.; Tulloch, J.; Henstridge, C.; Smith, C.; Spines-Jones, T.L. Clusterin accumulates in synapses in Alzheimer’s disease and is increased in apolipoprotein E4 carriers. Brain Commun. 2019, 1, fcz003. [CrossRef]

197. Galli, S.; Lopes, D.M.; Ammari, R.; Kopra, J.; Millar, S.E.; Gibb, A.; Salinas, P.C. Deficient Wnt signalling triggers striatal synaptic degeneration and impaired motor behaviour in adult mice. Nat. Commun. 2014, 5, 4992. [CrossRef]

198. Mao, B.; Wu, W.; Li, Y.; Hoppe, D.; Stannek, P.; Glinka, A.; Niehrs, C. LDL-receptor-related protein 6 is a receptor for Dickkopf proteins. Nat. Cell Biol. 2001, 411, 321–325. [CrossRef]

199. Bafico, A.; Liu, G.; Yaniv, A.; Gazit, A.; Aaronson, S.A. Novel mechanism of Wnt signalling inhibition mediated by Dickkopf-1 interaction with LRP6/Arrow. Nat. Cell Biol. 2001, 3, 683–686. [CrossRef]

200. Ross, S.P.; Baker, K.E.; Fisher, A.; Hoff, L.; Pak, E.S.; Muralov, A.K. miRNA-431 Prevents Amyloid-β-Induced Synapse Loss in Neuronal Cell Culture Model of Alzheimer’s Disease by Silencing Kremen1. Front. Cell. Neurosci. 2018, 12, 87. [CrossRef] [PubMed]

201. Sellers, K.J.; Elliott, C.; Jackson, J.; Ghosh, A.; Ribe, E.; Rojo, A.I.; Jarosz-Griffiths, H.H.; Watson, I.A.; Xia, W.; Semenov, M.; et al. Amyloid β synaptotoxicity is Wnt-PCP dependent and blocked by fasudil. Alzheimer’s Dement. 2018, 14, 306–317. [CrossRef] [PubMed]

202. Kettenmann, H.; Hanisch, U.-K.; Noda, M.; Verkhratsky, A. Physiology of Microglia. Physiol. Rev. 2011, 91, 461–553. [CrossRef] [PubMed]

203. Siskova, Z.; Tremblay, M.-È. Microglia and Synapse: Interactions in Health and Neurodegeneration. Neural Plast. 2013, 2013, 425845. [CrossRef] [PubMed]
204. Bradshaw, E.M.; Initiative, T.A.D.N.; Chibnik, L.B.; Keenan, B.T.; Ottoboni, L.; Raj, T.; Tang, A.; Rosenkrantz, L.L.; Imboywa, S.; Lee, M.; et al. CD33 Alzheimer’s disease locus: Altered monocyte function and amyloid biology. *Nat. Neurosci.* 2013, 16, 848–850. [CrossRef] [PubMed]

205. Heneka, M.T.; Carson, M.J.; El Khoury, J.; Landreth, G.E.; Brosseron, F.; Feinstein, D.L.; Jacobs, A.H.; Wyss-Coray, T.; Vitorica, J.; Ransohoff, R.M.; et al. Neuroinflammation in Alzheimer’s disease. *Lancet Neurol.* 2015, 14, 388–405. [CrossRef]

206. Henstridge, C.M.; Hyman, B.T.; Spires-Jones, T.L. Beyond the neuron–cellular interactions early in Alzheimer disease pathology. *Nat. Rev. Neurosci.* 2019, 20, 94–108. [CrossRef] [PubMed]

207. Keren-Shaul, H.; Spinrad, A.; Weiner, A.; Matcovitch-Natan, O.; Dvir-Szternfeld, R.; Ulland, T.K.; David, E.; Baruch, K.; Lara-Astaiso, D.; Toth, B.; et al. A Unique Microglia Type Associated with Restricting Development of Alzheimer’s Disease. *Cell* 2017, 169, 1276–1290.e17. [CrossRef]

208. Frigerio, C.S.; Wolfs, L.; Fattorelli, N.; Thrupp, N.; Voytyuk, I.; Schmidt, I.; Mancuso, R.; Chen, W.-T.; Bradshaw, E.M.; Initiative, T.A.D.N.; Chibnik, L.B.; Keenan, B.T.; Ottoboni, L.; Raj, T.; Tang, A.; Bamberger, M.E.; Harris, M.E.; McDonald, D.R.; Husemann, J.; Landreth, G.E. A Cell Surface Receptor Complex for Fibrillar β-Amyloid Mediates Microglial Activation. *J. Neurosci.* 2003, 23, 2665–2674. [CrossRef]

209. Meilandt, W.J.; Ngu, H.; Gogineni, A.; Lalehzadeh, G.; Lee, S.-H.; Srinivasan, K.; Imperio, J.; Wu, T.; Weber, M.; Keren-Shaul, H.; Spinrad, A.; Weiner, A.; Matcovitch-Natan, O.; Dvir-Szternfeld, R.; Ulland, T.K.; David, E.; Baruch, K.; Lara-Astaiso, D.; Toth, B.; et al. A Unique Microglia Type Associated with Restricting Development of Alzheimer’s Disease. *Cell* 2017, 169, 1276–1290.e17. [CrossRef]

210. Wang, Y.; Ulland, T.K.; Ulrich, J.D.; Song, W.; Tzaferis, J.A.; Hole, J.T.; Yuan, P.; Mahan, T.E.; Shi, Y.; Woodbury, M.E.; Srivastava, G.; et al. TREM2-mediated early microglial response limits diffusion and toxicity of amyloid plaques. *J. Exp. Med.* 2016, 213, 667–675. [CrossRef]

211. Paresce, D.M.; Ghosh, R.N.; Maxfield, F.R. Microglial Cells Internalize Aggregates of the Alzheimer’s Disease β-Protein Via a Scavenger Receptor. *Neuron* 1996, 17, 553–565. [CrossRef]

212. Bamberger, M.E.; Harris, M.E.; McDonald, D.R.; Husemann, J.; Landreth, G.E. A Cell Surface Receptor Complex for Fibrillar β-Amyloid Mediates Microglial Activation. *J. Neurosci.* 2003, 23, 2665–2674. [CrossRef]

213. Zhao, Y.; Wu, X.; Li, X.; Jiang, L.-L.; Gui, X.; Liu, Y.; Sun, Y.; Zhu, B.; Piña-Crespo, J.C.; Zhang, M.; et al. TREM2 Is a Receptor for β-Amyloid that Mediates Microglial Function. *Neuron* 2018, 97, 1023–1031.e7. [CrossRef]

214. Okun, E.; Griffioen, K.J.; Lathia, J.D.; Tang, S.-C.; Mattson, M.P.; Arumugam, T.V. Toll-like receptors in neurodegeneration. *Brain Res. Rev.* 2009, 59, 278–292. [CrossRef] [PubMed]

215. Condello, C.; Yuan, P.; Schain, A.; Grutzendler, J. Microglia constitute a barrier that prevents neurotoxic protofibrillar Aβ42 hotspots around plaques. *Nat. Commun.* 2015, 6, 7176. [CrossRef]

216. Spangenberg, E.E.; Lee, R.J.; Najafi, A.R.; Rice, R.A.; Elmore, M.R.P.; Blunting-Jones, M.; West, B.L.; Green, K.N. Eliminating microglia in Alzheimer’s mice prevents neuronal loss without modulating amyloid-β pathology. *Brain* 2016, 139, 1265–1281. [CrossRef] [PubMed]

217. Grathwohl, S.A.; Kalin, R.E.; Bolmont, T.; Prokop, S.; Winkelmann, G.; Kaeser, S.A.; Odenthal, J.; Radde, R.; Eldh, T.; Gandy, S.; et al. Formation and maintenance of Alzheimer’s disease β-amyloid plaques in the absence of microglia. *Nat. Neurosci.* 2009, 12, 1361–1363. [CrossRef] [PubMed]

218. Zhao, R.; Hu, W.; Tsai, J.; Li, W.; Gan, W.-B. Microglia limit the expansion of β-amyloid plaques in a mouse model of Alzheimer’s disease. *Mol. Neurodegener.* 2017, 12, 1–11. [CrossRef]

219. Cziriz, E.; Castello, N.A.; Mosher, K.I.; Castellano, J.M.; Hinkson, I.V.; Lucin, K.M.; Baeza-Raja, B.; Ryu, J.K.; Li, L.; Farina, S.N.; et al. Microglial complement receptor 3 regulates brain Aβ levels through secreted proteolytic activity. *J. Exp. Med.* 2017, 214, 1081–1092. [CrossRef]

220. Akiyama, H.; Barger, S.; Barnum, S.; Bradt, B.; Bauer, J.; Cole, G.M.; Cooper, N.R.; Emmerling, M.; Fiebich, B.L.; et al. Inflammation and Alzheimer’s disease. *Neurobiol. Aging* 2000, 21, 383–421. [CrossRef]

221. Hong, S.; Beja-Glasser, V.F.; Nfonoyim, B.M.; Frouin, A.; Li, S.; Ramakrishnan, S.; Merry, K.M.; Shi, Q.; Rosenthal, A.; Barres, B.A.; et al. Complement and microglia mediate early synapse loss in Alzheimer mouse models. *Science* 2016, 352, 712–716. [CrossRef]

222. Rajendran, L.; Paolicelli, R.C. Microglia-Mediated Synapse Loss in Alzheimer’s Disease. *J. Neurosci.* 2018, 38, 2911–2919. [CrossRef]

223. Ma, B.; Hottiger, M.O. Crosstalk between Wnt/β-Catenin and NF-kB Signaling Pathway during Inflammation. *Front. Immunol.* 2016, 7, 378. [CrossRef]
Brain Sci. 2020, 10, 902

224. Zolezzi, J.M.; Inestrosa, N.C. Wnt/TLR Dialog in Neuroinflammation, Relevance in Alzheimer’s Disease. *Front. Immunol.* 2017, *8*, 187. [CrossRef] [PubMed]

225. Liu, S.; Liu, Y.; Hao, W.; Wolf, L.; Kiliaan, A.J.; Penke, B.; Rübe, C.E.; Walter, J.; Heneka, M.T.; Hartmann, T.; et al. TLR2 Is a Primary Receptor for Alzheimer’s Amyloid β Peptide To Trigger Neuroinflammatory Activation. *J. Immunol.* 2011, *188*, 1098–1107. [CrossRef]

226. Reed-Geaghan, E.G.; Savage, J.C.; Hise, A.G.; Landreth, G.E. CD14 and Toll-Like Receptors 2 and 4 Are Required for Fibrillar Aβ-Stimulated Microglial Activation. *J. Neurosci.* 2009, *29*, 11982–11992. [CrossRef] [PubMed]

227. Ma, B.; Fey, M.; Hottiger, M.O. WNT/β-catenin signaling inhibits CBP-mediated RelA acetylation and expression of proinflammatory NF-κB target genes. *J. Cell Sci.* 2015, *128*, 2430–2436. [CrossRef] [PubMed]

228. Noubissi, F.K.; Elcheva, I.; Bhatia, N.; Shakoori, A.; Ougolkov, A.; Liu, J.; Minamoto, T.; Ross, J.; Fuchs, S.Y.; Spiegelman, V.S. CRD-BP mediates stabilization of βTrCP1 and c-myc mRNA in response to β-catenin signalling. *Nat. Cell Biol.* 2006, *441*, 898–901. [CrossRef] [PubMed]

229. Van Steenwinckel, J.; Schang, A.-L.; Krishnan, M.L.; Degos, V.; Delahaye-Duriez, A.; Bokobza, C.; Csaba, Z.; Verdonk, F.; Montané, A.; Sigaut, S.; et al. Decreased microglial Wnt/β-catenin signalling drives microglial pro-inflammatory activation in the developing brain. *Brain* 2019, *142*, 3806–3833. [CrossRef]

230. Zhang, D.; Lu, Z.; Man, J.; Cui, K.; Fu, X.; Yu, L.; Gao, Y.; Liao, L.; Xiao, Q.; Guo, R.; et al. Wnt-3a alleviates neuroinflammation after ischemic stroke by modulating the responses of microglia/macrophages and astrocytes. *Int. Immunopharmacol.* 2019, *75*, 105760. [CrossRef]

231. Stamatovic, S.M.; Johnson, A.M.; Keep, R.F.; Andjelkovic, A.V. Junctional proteins of the blood-brain barrier: A Standardized Model of the Blood Brain Barrier In Vitro. *PLoS ONE* 2013, *8*, e70233. [CrossRef] [PubMed]
Brain Sci. 2020, 10, 902

245. Hübner, K.; Caboche, P.; Diéguez-Hurtado, R.; Wiesner, C.; Wakayama, Y.; Grassme, K.S.; Hubert, M.; Guenther, S.; Belting, H.-G.; Affolter, M.; et al. Wnt/β-catenin signaling regulates VE-cadherin-mediated anastomosis of brain capillaries by counteracting S1pr1 signaling. Nat. Commun. 2018, 9, 4860. [CrossRef] [PubMed]

246. Lim, J.C.; Kania, K.D.; Wijesuriya, H.; Chawla, S.; Sethi, J.K.; Pulaski, L.; Romero, I.A.; Couraud, P.O.; Weksler, B.B.; Hladky, S.B.; et al. Activation of β-catenin signalling by GSK-3 inhibition increases P-glycoprotein expression in brain endothelial cells. J. Neurochem. 2008, 106, 1855–1865. [CrossRef] [PubMed]

247. Laksitorini, M.D.; Yathindranath, V.; Xiong, W.; Hombach-Klonisch, S.; Miller, D.W. Modulation of Wnt/β-catenin signaling promotes blood-brain barrier phenotype in cultured brain endothelial cells. Sci. Rep. 2019, 9, 4860. [CrossRef] [PubMed]

248. Pinzón-Daza, M.L.; Salaroglio, I.C.; Kopecka, J.; Garzón, R.; Couraud, P.-O.; Ghigo, D.; Riganti, C. The Cross-Talk between Canonical and Non-Canonical Wnt-Dependent Pathways Regulates P-Glycoprotein Expression in Human Blood–Brain Barrier Cells. Br. J. Pharmacol. 2014, 34, 1258–1269. [CrossRef] [PubMed]

249. Tran, K.A.; Zhang, X.; Predescu, D.; Huang, X.; Machado, R.F.; Göthert, J.R.; Malik, A.B.; Valyi-Nagy, T.; Zhao, Y.-Y. Endothelial β-Catenin Signaling Is Required for Maintaining Adult Blood–Brain Barrier Integrity and Central Nervous System Homeostasis. Circulation 2016, 133, 177–186. [CrossRef]

250. Wang, Y.; Cho, C.; Williams, J.; Smallwood, P.M.; Zhang, C.; Junge, H.J.; Nathans, J. Interplay of the Norrin and Wnt7a/Wnt7b signaling systems in blood–brain barrier and blood–retina barrier development and maintenance. Proc. Natl. Acad. Sci. USA 2018, 115, E11827–E11836. [CrossRef] [PubMed]

251. Kania, K.D.; Wijesuriya, H.C.; Hladky, S.B.; Barrand, M.A. Beta amyloid effects on expression of multidrug efflux transporters in brain endothelial cells. Brain Res. 2011, 1418, 1–11. [CrossRef]

252. Wijesuriya, H.C.; Bullock, J.Y.; Faull, R.L.; Hladky, S.B.; Barrand, M.A. ABC efflux transporters in brain vasculature of Alzheimer’s subjects. Brain Res. 2010, 1358, 228–238. [CrossRef]

253. Lim, J.C.; Mickute, Z.; Zaman, M.; Hopkins, S.; Wijesuriya, H.; Steckerl, T.; Moechars, D.; Van Leuven, F.; Sarnyai, Z.; Hladky, S.B.; et al. Decreased expression of multidrug efflux transporters in the brains of GSK-3β transgenic mice. Brain Res. 2009, 1276, 1–10. [CrossRef]

254. Cirrito, J.R.; Deane, R.; Fagan, A.M.; Spinner, M.L.; Parsadanian, M.; Finn, M.B.; Jiang, H.; Prior, J.L.; Sagare, A.; Bales, K.R.; et al. P-glycoprotein deficiency at the blood-brain barrier increases amyloid- deposition in an Alzheimer disease mouse model. J. Clin. Investig. 2005, 115, 3285–3290. [CrossRef] [PubMed]

255. Miller, M.C.; Tavares, R.; Johanson, C.E.; Hovanesian, V.; Donahue, J.E.; Gonzalez, L.; Silverberg, G.D.; Stopa, E.G. Hippocampal RAGE immunoreactivity in early and advanced Alzheimer’s disease. Brain Res. 2008, 1230, 273–280. [CrossRef] [PubMed]

256. Hartz, A.M.S.; Miller, D.S.; Bauer, B. Restoring blood-brain barrier P-glycoprotein reduces brain amyloid-beta in a mouse model of Alzheimer’s disease. Mol. Pharmacol. 2010, 77, 715–723. [CrossRef] [PubMed]

257. Mooradian, A.; Chung, H.; Shah, G. GLUT-1 Expression in the Cerebria of Patients with Alzheimer’s Disease. Neurobiol. Aging 1997, 18, 469–474. [CrossRef]

258. Winkler, E.A.; Nishida, Y.; Sagare, A.P.; Rege, S.V.; Bell, R.D.; Perlmutter, D.; Sengillo, J.D.; Hillman, S.; Kong, P.; Nelson, A.R.; et al. GLUT1 reductions exacerbate Alzheimer’s disease vasculo-neuronal dysfunction and degeneration. Nat. Neurosci. 2015, 18, 521–530. [CrossRef] [PubMed]

259. Hou, Y.; Dan, X.; Babbar, M.; Wei, Y.; Hasselbalch, S.G.; Croteau, D.L.; Bohr, V.A. Ageing as a risk factor for neurodegenerative disease. Nat. Rev. Neuro. 2019, 15, 565–581. [CrossRef]

260. Mu, Q.; Chen, Y.; Wang, J. Deciphering Brain Complexity Using Single-cell Sequencing. Genom. Proteom. Bioinfom. 2019, 17, 344–366. [CrossRef]

261. Tay, L.; Leung, B.; Yeo, A.; Chan, M.; Lim, W.S. Elevations in Serum Dickkopf-1 and Disease Progression in Community-Dwelling Older Adults With Mild Cognitive Impairment and Mild-to-Moderate Alzheimer’s Disease. Front. Aging Neurosci. 2019, 11, 278. [CrossRef]

262. Zhu, Z.; Guo, D.; Zhong, C.; Wang, A.; Xie, X.; Xu, T.; Chen, C.-S.; Peng, Y.; Peng, H.; Li, Q.; et al. Serum Dkk-1 (Dickkopf-1) Is a Potential Biomarker in the Prediction of Clinical Outcomes Among Patients With Acute Ischemic Stroke. Arter. Thromb. Vasc. Biol. 2019, 39, 285–293. [CrossRef]

263. Nusse, R.; Clevers, H. Wnt/β-Catenin Signaling, Disease, and Emerging Therapeutic Modalities. Cell 2017, 169, 985–999. [CrossRef]
264. Behl, T.; Kaur, I.; Fratila, O.; Brata, R.; Bungau, S. Exploring the Potential of Therapeutic Agents Targeted towards Mitigating the Events Associated with Amyloid-β Cascade in Alzheimer’s Disease. *Int. J. Mol. Sci.* **2020**, *21*, 7443. [CrossRef] [PubMed]

265. Vaz, M.; Silvestre, S. Alzheimer’s disease: Recent treatment strategies. *Eur. J. Pharmacol.* **2020**, *887*, 173554. [CrossRef] [PubMed]

266. Hosseinpour, S.; Rad, M.R.; Khojasteh, A.; Zadeh, H. Antibody Administration for Bone Tissue Engineering: A Systematic Review. *Curr. Stem Cell Res. Ther.* **2018**, *13*, 292–315. [CrossRef] [PubMed]

267. Li, B.; Rong, D.; Wang, Y. Targeting Protein-Protein Interaction with Covalent Small-Molecule Inhibitors. *Curr. Top. Med. Chem.* **2019**, *19*, 1872–1876. [CrossRef] [PubMed]

268. Silva, A.C.; Lobo, D.D.; Martins, I.M.; Lopes, S.M.; Henriques, C.; Duarte, S.P.; Dodart, J.-C.; Nobre, R.; De Almeida, L.P. Antisense oligonucleotide therapeutics in neurodegenerative diseases: The case of polyglutamine disorders. *Brain* **2019**, *143*, 407–429. [CrossRef]

269. Lewcock, J.W.; Schlepckow, K.; Di Paolo, G.; Tahirovic, S.; Monroe, K.M.; Haass, C. Emerging Microglia Biology Defines Novel Therapeutic Approaches for Alzheimer’s Disease. *Neuron* **2020**, *6273*, 30753–30754. [CrossRef]

270. Lengfeld, J.E.; Lutz, S.E.; Smith, J.R.; Diaconu, C.; Scott, C.; Kofman, S.B.; Choi, C.; Walsh, C.M.; Raine, C.S.; Agalliu, I.; et al. Endothelial Wnt/β-catenin signaling reduces immune cell infiltration in multiple sclerosis. *Proc. Natl. Acad. Sci. USA* **2017**, *114*, E1168–E1177. [CrossRef] [PubMed]

271. Leblanc, N.J.; Menet, R.; Picard, K.; Parent, G.; Tremblay, M.-È.; ElAli, A. Canonical Wnt Pathway Maintains Blood-Brain Barrier Integrity upon Ischemic Stroke and Its Activation Ameliorates Tissue Plasminogen Activator Therapy. *Mol. Neurobiol.* **2019**, *56*, 6521–6538. [CrossRef]

272. Salehi, A.; Jullienne, A.; Baghchechi, M.; Hamer, M.; Walsworth, M.; Donovan, V.; Tang, J.; Zhang, J.H.; Pearce, W.J.; Obenaus, A. Up-regulation of Wnt/β-catenin expression is accompanied with vascular repair after traumatic brain injury. *Br. J. Pharmacol.* **2017**, *38*, 274–289. [CrossRef]

273. Wold, E.A.; Chen, J.; Cunningham, K.A.; Zhou, J. Allosteric Modulation of Class A GPCRs: Targets, Agents, and Emerging Concepts. *J. Med. Chem.* **2019**, *62*, 88–127. [CrossRef]

274. Molinuevo, J.L.; Ayton, S.; Battrla, R.; Bednar, M.M.; Bittner, T.; Cummings, J.; Fagan, A.M.; Hampel, H.; Mielke, M.M.; Mikulskis, A.; et al. Current state of Alzheimer’s fluid biomarkers. *Acta Neuropathol.* **2018**, *136*, 821–853. [CrossRef] [PubMed]

275. Park, S.A.; Han, S.M.; Kim, C.E. New fluid biomarkers tracking non-amyloid-β and non-tau pathology in Alzheimer’s disease. *Exp. Mol. Med.* **2020**, *52*, 556–568. [CrossRef] [PubMed]

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