ABSTRACT

The treatment of patients with early breast cancer has always been characterised by escalation by new therapies and de-escalation through identification of better treatment regimens or introduction of better tools to estimate prognosis. Efforts in some of these areas in the last few years have led to solid data. The results of the large studies of de-escalation through use of multi-gene tests are available, as are the results of some studies that investigated the new anti-HER2 substances.
T-DM1 and pertuzumab in the early treatment situation. Several large-scale studies examining the role of CDK4/6 inhibitors will soon be concluded so innovations can be anticipated in this area also. This review article will summarise and classify the results of the latest publications.

ZUSAMMENFASSUNG

Die Therapie von Patientinnen mit frühem Mammakarzinom war immer schon geprägt von Eskalation durch neue Therapien und Deeskalation durch Identifikation besserer Therapieschemata oder Einführung von besseren Werkzeugen zur Einschätzung der Prognose. Die Anstrengungen der letzten Jahre haben auf einigen dieser Gebiete zu einer soliden Datenlage geführt. Die Ergebnisse der großen Deeskalationstudien durch Nutzung von Multi-Gen-Tests liegen ebenso vor wie die Ergebnisse einiger Studien, welche die neuen Anti-HER2-Substanzen T-DM1 und Pertuzumab in der frühen Therapiesituation untersucht haben. Mehrere groß angelegte Studien zur Untersuchung der Rolle der CDK4/6-Inhibitoren stehen kurz vor dem Abschluss, sodass auch in dieser Therapiesituation mit Neuerungen zu rechnen ist. Diese Übersichtsarbeit soll die Ergebnisse der neuesten Publikationen zusammenfassen und einordnen.

Introduction

The treatment of patients with early breast cancer has changed in recent years, especially of patients with HER2-positive tumours through the introduction of T-DM1 and pertuzumab. CDK4/6 inhibitors for HER2-negative, hormone receptor-positive tumours could also be added soon, though the patient population is unclear since one of the adjuvant studies announced a negative study outcome (PALLAS) according to the press release and another study announced a positive result, likewise by press release (MonarchE). Another study in this indication is still recruiting (NataLEE). Until further major changes are possible implemented clinically in the (neo-)adjuvant situation, there have in the meantime been other interesting insights into the mode of action of existing therapies for many clinical scenarios. This review article will summarise the recent publications from international conferences.

Prevention

Use of knowledge of risk factors for prevention

Epidemiological studies and the recording of genetic and other risk factors are becoming increasingly detailed and comprehensive so that a relatively good estimate of the magnitude of the individual breast cancer risk can be made. One in eight women will develop breast cancer up to the age of 85 years. Even though mortality is decreasing because of improved early diagnosis and treatment, the incidence of breast cancer has not fallen but has even increased in Western industrialised countries. With all the scientific efforts of recent decades, the question arises of how the knowledge can be used actually to reduce the incidence of breast cancer (primary prevention).

Among the genetic risk factors, a distinction is currently made between high-penetrance, moderate-penetrance and low-penetrance genetic changes. Most high-penetrance and moderate-penetrance genes are already being investigated today in panel gene tests as part of predictive genetic diagnostics. In addition to BRCA1 and BRCA2, which are still the most important for planning individual prevention, other genes such as PALB2, CHEK2, ATM and others have been genotyped [1–8]. Since these gene changes are present very rarely in the general population, however, it is difficult to envisage that broad genotyping of these genes can contribute to a reduction in disease rates.

In addition to the high- and moderate-penetrance genes, further risk variants have been identified in over 150 genomic regions [9–26]. Although these low-penetrance risk variants occur relatively frequently in the population, they have only a slight effect individually on the individual breast cancer risk. All genetic risk variants together explain roughly 40% of the increased familial breast cancer risk.

Among the non-genetic risk factors, mammographic density has the greatest effect on breast cancer risk. High mammographic density (> 50%) is present in ca. 20% of women. These have an approximately three-fold increased incidence of breast cancer [27, 28]. Similarly to some genetic changes, this risk is not the same for all molecular subtypes [29, 30]. Mammographic density is also correlated with several genetic and non-genetic risk factors [29–36]. The few relevant protective factors include an early first childbirth, prolonged breast-feeding and possibly sports [37].

As mentioned above for genetic risk factors, the risk factors that have a large effect on disease risk occur rather rarely and the risk factors that have a small effect occur frequently in the population. This means that a marked increase in risk applies for only a few individuals in a population (Fig. 1). There are several models that attempt to integrate the genetic and non-genetic risk factors in risk models that can better quantify the individual risk. However, these have not yet been integrated in studies or treatment concepts [34, 36, 38–40].

Neoadjuvant Therapy

Monitoring of neoadjuvant therapy in patients with HER2-positive breast cancer

Patients with HER2-positive breast cancer are among the patients in whom pathological complete remission (pCR) correlates very strongly with a good prognosis after neoadjuvant therapy [41–43]. Against this background there is great interest in identifying patients early during neoadjuvant therapy in order to continue de-escalated therapy until surgery if appropriate. The recently reported PHERGAIN study was conducted in this context [44]. The study design is complex and shown in Fig. 2. The HER2-positive patients received either a standard treatment with 6 cycles of tax-
**Fig. 1** Risk relationship and frequency of different factors that increase or decrease the risk of breast cancer.

**Key eligibility criteria**

1. Centrally confirmed HER2\(^{+}\) stage I–IIIA EBC.
2. Tumor diameter ≥ 1.5 cm by MRI or ultrasound.
3. Presence of a breast PET-evaluable lesion.

**Stratification factors**

- Hormonal receptor status (+/−)

**After cycle 2 (6 weeks)**

- Arm A: TCHP × 2
- Arm B: PH (ETx) × 2
- Arm C: TCHP × 6

**After cycle 6 or 8**

- Response: PH (ETx) × 6
- No response: TCHP × 6

**First primary endpoint:**
- pCR in PET responders (Arm B)

**Second primary endpoint:**
- 3-year iDFS rate in Arm B

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**Fig. 2** PHERGAIN study design. C: Carboplatin; D: Docetaxel; EVC: Early breast cancer; ETx: Endocrine therapy (Letrozole post-menopausal/ Tamoxifen pre-menopausal) Adjuvant ETx up to 3 years from surgery; PET: \(^{18}\)F-fluorodeoxyglucose positron emission tomography/computed tomography; H: Trastuzumab SC; HER2: Human Epidermal Growth Factor Receptor 2; IDFS: Invasive disease-free survival; MRI: Magnetic resonance Imaging; P: Pertuzumab IV; R: Randomization; TCHIP: Trastuzumab, Pertuzumab, Docetaxel, and Carboplatin. \(^{1}\)All hormonal receptor-positive patients will receive ETx concomitantly with PH (except on chemotherapy) PET responders: RECIST responders after cycle 2 with SUV\(_{\text{max}}\) reduction ≥ 40%. pCR: Pathological complete response (ypT0/isN0). Modified after [44].
The use of endocrine-based therapy in the neoadjuvant situation represents an alternative to chemotherapy for a certain patient population and is currently being investigated intensively in studies [51–53]. In particular, a neoadjuvant study can investigate...
how certain resistance mechanisms are overcome by CDK4/6 inhibitors. Data regarding abemaciclib were already available from the Neo-Monarch study that showed that abemaciclib leads to marked cell cycle arrest in the neoadjuvant setting [54]. In this connection, the FELINE study, which investigated neoadjuvant use of ribociclib, has now been reported [55]. Patients with primary HER2-negative, HR-positive breast cancer were randomised to 3 treatment arms, each lasting for 6 months:
1. Letrozole monotherapy,
2. Letrozole + continuous ribociclib,
3. Letrozole + intermittently paused ribociclib.

The primary study aim was the frequency of a PEPI score of zero (0) after the neoadjuvant therapy [56]. Interestingly, the frequency of patients with a PEPI score of 0 did not differ between the letrozole monotherapy and the CDK4/6 therapy arms. The percentage was 25.8% in the letrozole monotherapy arm and 25.4% in the two ribociclib arms together (p = 0.96). It was shown, however, that complete cell cycle arrest was attained in 91.9% of patients treated with ribociclib after 14 days of treatment, while this was detectable in only 51.7% of the patients on letrozole monotherapy (p = 0.0001). This difference was smaller by the time of surgery (71.4% after 6 months of therapy containing ribociclib and 61.3% after 6 months of letrozole monotherapy, p = 0.4225). The FELINE study thus delivers interesting insights into how cell cycle arrest behaves when endocrine monotherapy is compared with CDK4/6 inhibitor therapy + ET.

Locoregional Therapies

Surgery of the primary tumour as part of primary treatment even if M1 at initial diagnosis?

Roughly 6–10% of patients newly diagnosed with breast cancer already have distant metastases at the time of diagnosis. For these patients the question arises as to whether surgery of the local disease should be performed as part of the initial treatment. A few retrospective studies had implied this, but the analyses were not balanced. Patients who had had surgery were generally younger, had smaller tumours, and had more often had hormone receptor-positive disease and less advanced malignant disease [57]. Two prospective randomised studies yielded conflicting results [58, 59]. In this connection the new E2108 study has been reported [57]. This study randomised 256 patients who had shown no progression with primary systemic therapy. 131 of these patients continued to receive the systemic therapy and 125 patients had surgery after initial systemic therapy. The overall survival, which was the primary study aim, did not differ between the two randomisation arms. The hazard ratio was 1.09 (90% CI: 0.80–1.49). There was no difference in progression-free survival either. With regard to locoregional recurrence, this occurred in 10.2% of cases in the surgery arm and locoregional recurrence or progression occurred in 5 and 20.6% of cases in the randomisation arms without surgery. However, this did not affect quality of life. The authors of the study concluded that when the disease is well controlled by systemic therapy, surgery could take place only when the local disease progresses.

Adjuvant Therapy

T-DM1 to avoid adjuvant chemotherapy?

The antibody-drug conjugate (ADC) T-DM1 is in clinical use both in patients with advanced HER2-positive breast cancer and in the post-neoadjuvant situation [60,61]. This naturally raised the question of whether T-DM1, possibly in combination with pertuzumab, could replace a therapy that includes conventional chemotherapy. KRISTINE/TRIO-021 is a study in this connection that has already been conducted in the neoadjuvant setting. In this neoadjuvant study, treatment with TCHP was compared to treatment with T-DM1 + pertuzumab. Treatment with TCHP led to a significantly higher pCR (56%) compared with 44% in the T-DM1 + pertuzumab arm [62]. With regard to the prognosis, more events were apparent in the T-DM1 + pertuzumab arm than in the TCHP arm, most probably due to preoperative progression [63].

In this connection, the KAITLIN study has now investigated the combination of T-DM1 + pertuzumab in the adjuvant situation also [64]. The KAITLIN study compared treatment with AC-THP with treatment with T-DM1 + pertuzumab in a mainly node-positive HER2-positive population of primary breast cancer patients after surgery. 1846 patients were included in the study. The invasive recurrence-free survival (primary study aim) did not differ between the two treatments. The hazard ratio was 0.97 (95% CI: 0.71–1.32). It must be commented that the invasive recurrence-free 3-year survival was very good for the high-risk patients included in the study, at 94.1% in the AC-THP arm and 92.8% in the AC-DM1/P arm. Grade 3/4 adverse effects were similar in the two arms. This rate was 55.4% in the AC-THP arm and 51.8% of the patients in the AC-DM1/P arm suffered a grade 3/4 adverse effect. The authors concluded that treatment with AC-THP continues to be the standard treatment for patients with HER2-positive breast cancer.

Consolidated data from the MINDACT study for decision-making regarding adjuvant therapy

Besides the TailorX study, which investigated a 21-gene score with regard to decision-making in HER2-negative, hormone receptor-positive patients [65,66], the MINDACT study is the second large study that addresses the question of whether and which patients in this population can be spared chemotherapy. The MINDACT study bases its genomic analyses on a 70-gene risk score [67]. The study design is shown in ▶ Figs. 4 and 5 shows a comparison of the TailorX and MINDACT study populations. The primary aim of the MINDACT study was reached [67], which was defined that patients with a high clinical recurrence risk and who had not received any chemotherapy based on the genomic assessment should have a better metastasis-free 5-year survival than 92%. The metastasis-free 5-year survival at the time was 94.7% (95% CI: 92.5–96.2%) [67]. Long-term follow-up observations after 8.7 years have now been presented. The primary analysis was confirmed with the more mature data. The metastasis-free 5-year survival was 95.1% (95% CI: 93.1–96.6%) [68].

Consideration of the four groups who were observed in the MINDACT study confirmed the results of the primary analysis
No additional therapy:
- 78.1 alive in 10 years
- 16.6 die of cancer
- 5.3 die of other causes

With hormonal therapy:
Benefit = 4.9 alive

With chemotherapy:
Benefit = 2.4 alive

With combined therapy:
Benefit = 6.6 alive

Registration and screening surgery
n = 6693

Clinical-pathological (C) risk
Genomic (G) risk
(70-gene signature)
Discordant cases
C-low/G-high or C-high/G-low
1st randomization to treatment
use clinical vs. genomic risk

2nd randomization
Anthracycline-based vs. Capecitabine-Docetaxel

Endocrine therapy

3rd randomization
Tamoxifen 2 years/Letrozole 5 years vs. Letrozole 7 years

MINDACT population:
HR+/HER2− 81%
HER2+ 9.5%
TNBC 9.6%

▶ Fig. 4 MINDACT study design. HER2: Human Epidermal Growth Factor Receptor 2; HR: hormone receptor; TNBC: triple negative breast cancer. Modified after [68].

MINDACT population clinical high/MammaPrint low
n = 1551 (577 premenopausal)Median age = 55 years
Node (1–3) positive 48%
T-size > 2 cm 58%
Grade 3 29%
Clinical high risk 100%

TailorX population recurrence score 11–26
n = 6711 (2415 premenopausal)Median age = 55 years
Node (1–3) positive None
T-size > 2 cm 24%
Grade 3 14%
Clinical high risk 26%
Clinical low risk 74%

In HR+/HER2− C-high/G-low patients: 49% node (1–3) positive and 27% grade 3

▶ Fig. 5 Comparison of the TailorX and MammaPrint study populations. HER2: Human Epidermal Growth Factor Receptor 2; HR: hormone receptor; TN: triple negative. Modified after [68].
Patients who have a good prognosis according to both assessment methods (clinical and genomic) have an excellent prognosis without chemotherapy. Patients who have a poor prognosis as assessed clinically and genomically had a poor prognosis even despite chemotherapy. Both groups with a discordant estimate of prognosis had a similar prognosis; chemotherapy can therefore be avoided for this group.

Biomarkers

T-DM1 to avoid adjuvant chemotherapy?

The introduction of T-DM1 after neoadjuvant anti-HER2 therapy without pCR has signified a marked improvement in treatment results for these patients [61]. The study that delivered these results (KATHERINE) integrated a very comprehensive translational research programme prospectively in the study procedure. This offers the possibility of investigating the resistance mechanisms of different anti-HER2 therapies in order to identify patients who may react particularly well to adjuvant therapy with T-DM1 after a suboptimal response to neoadjuvant therapy. An analysis has now been presented that examined tumour samples before and after the neoadjuvant therapy. This focused on the PI3K signalling pathway and the gene expression profiles of the immune signalling pathways [69]. Whether PIK3CA mutations have an influence on the prognosis of the patients in the study was to be investi-

gated. In the overall population, no influence was seen on invasive recurrence-free survival. The hazard ratios were similar in all groups (Fig. 7) and were 0.48 (95% CI: 0.35–0.65) in patients with non-mutated tumours and 0.54 (95% CI: 0.32–0.90) in mutated patients [69]. Post-neoadjuvant therapy is therefore independent of the PIK3CA mutation status.

For the immunological analyses, samples were examined at the time of surgery with regard to the expression of HER2, PD-L1, CD8 and T cell effector molecules. None of these gene ex-

Fig. 6 Survival rates in the 4 groups of the MINDACT study. c: clinical; g: genomic; H: high; L: low. Modified after [68].

Fig. 7 Effect of T-DM1 therapy in the KATHERINE study according to PIK3CA mutation status. CI: confidence interval; HR: hazard ratio; IDFS: invasive disease-free survival; T-DM1: trastuzumab emtansine [69].
pression signatures was able to identify a group of patients in which T-DM1 had different effectiveness than in other subgroups.

It was interesting that high HER2 expression after neoadjuvant anti-HER2 therapy provided evidence for resistance for adjuvant trastuzumab. In patients with high HER2 expression after neoadjuvant therapy the risk for a recurrence event was twice as high compared with patients with low HER2 expression (hazard ratio: 2.02; 95% CI: 1.32–3.11). This was not the case in patients who had received adjuvant T-DM1 (hazard ratio: 1.01; 95% CI: 0.56–1.83) [69].

Outlook

Even though there are few data at present regarding an improvement of the treatment of patients with triple-negative cancer, an attempt is now being made to translate the results of studies in the advanced therapy situation to the treatment of early breast cancer. The ADC sacituzumab govitacan, which has shown clear efficacy in metastatic TNBC [70], is currently being integrated in a larger study of HER2-negative patients in the post-neoadjuvant setting (SASCIA study [71]). The results of the large adjuvant CDK4/6 inhibitor studies will also be interesting as there is a high probability that these will change routine clinical practice in the near future.

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Conflict of Interest

A. D. H. received speaker and consultancy honoraria from AstraZeneca, Genomic Health, Roche, Novartis, Celgene, Lilly, MSD, Eisai, Teva, Tesaro, Daiichi-Sankyo, Hexal and Pfizer. F. O. received speaker and consultancy honoraria from Amgen, AstraZeneca, Bayer, BMS, Boehringer-Ingelheim, Chugai, Celgene, Cellex, Eisai, Gilead, Hexal, Ipsen, Janssen-Cilag, Merck, MSD, Novartis, Novonordic, Riemsier, Roche, Servier, Shire, Tesaro, TEVA. H.-C. K. received honoraria from Carl Zeiss medic, Teva, Theracration, Novartis, Amgen, AstraZeneca, Pfizer, Janssen-Cilag, GSK, Liv Pharma, Roche and Genomic Health. P. A. F. received honoraria from Novartis, Pfizer, Roche, Amgen, Celgene, Daiichi Sankyo, AstraZeneca, Merck-Sharp & Dohme, Eisai, Puma and Teva. His institution conducts research with funding from Novartis and Biontech. H. T. received honoraria from Novartis, Roche, Celgene, Teva, Pfizer and travel support from Roche, Celgene and Pfizer. J. E. received honoraria from AstraZeneca, Roche, Celgene, Novartis, Lilly, Pfizer, Pierre Fabre, Teva and travel support from Celgene, Pfizer, Teva and Pierre Fabre.

M. P. L. has participated on advisory boards for AstraZeneca, Lilly, MSD, Novartis, Pfizer, Eisai, Genomic Health and Roche and has received honoraria for lectures from MSD, Roche, Novartis, Pfizer, Genomic Health, AstraZeneca, Celgene, Lilly, Loreal, MSD, Novartis, Pfizer, Tesaro, Teva T. N. F. has participated on advisory boards for Amgen, Daiichi Sankyo, Novartis, Pfizer, and Roche and has received honoraria for lectures from Amgen, Celgene, Daiichi Sankyo, Roche, Novartis and Pfizer. M. T. has participated on advisory boards for AstraZeneca, Lilly, MSD, Novartis, Pfizer, Genomic Health and Roche and has received honoraria for lectures from MSD, Roche, Novartis, Pfizer, Genomic Health, and AstraZeneca M.W. has participated on advisory boards for AstraZeneca, Lilly, MSD, Novartis, Pfizer and Roche. J. H. reports receiving speakers bureau honoraria from Celgene, Novartis, and Roche, and is a consultant/advisory board member for Amgen, Celgene, Novartis and Roche.

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