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To cite this version:
Sandrine Richard, Frédéric Selle, Jean-Pierre Lotz, Ahmed Khalil, Joseph Gligorov, et al.. Pertuzumab and trastuzumab: the rationale way to synergy. Anais da Academia Brasileira de Ciências, Academia Brasileira de Ciências, 2016, 88 (supl.1), 10.1590/0001-3765201620150178. hal-01334010

HAL Id: hal-01334010
https://hal.sorbonne-universite.fr/hal-01334010
Submitted on 20 Jun 2016

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Pertuzumab and trastuzumab: the rationale way to synergy

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Manuscript received on March 13, 2015; accepted for publication on May 5, 2015

ABSTRACT

It has now been 15 years since the HER2-targeted monoclonal antibody trastuzumab was introduced in clinical and revolutionized the treatment of HER2-positive breast cancer patients. Despite this achievement, most patients with HER2-positive metastatic breast cancer still show progression of their disease, highlighting the need for new therapies. The continuous interest in novel targeted agents led to the development of pertuzumab, the first in a new class of agents, the HER dimerization inhibitors. Pertuzumab is a novel recombinant humanized antibody directed against extracellular domain II of HER2 protein that is required for the heterodimerization of HER2 with other HER receptors, leading to the activation of downstream signalling pathways. Pertuzumab combined with trastuzumab plus docetaxel was approved for the first-line treatment of patients with HER2-positive metastatic breast cancer and is currently used as a standard of care in this indication. In the neoadjuvant setting, the drug was granted FDA-accelerated approval in 2013. Pertuzumab is also being evaluated in the adjuvant setting. The potential of pertuzumab relies in the dual complete blockade of the HER2/3 axis when administered with trastuzumab. This paper synthesizes preclinical and clinical data on pertuzumab and highlights the mechanisms underlying the synergistic activity of the combination pertuzumab-trastuzumab which are essentially due to their complementary mode of action.

Key words: breast cancer, dimerization, HER2/3, monoclonal antibody, pertuzumab, trastuzumab.

INTRODUCTION

The human epidermal growth factor receptor 2 (HER2, ErbB-2 or HER-2/neu) gene, which encodes the HER2 receptor tyrosine kinase, is amplified in about 20% of breast cancers (Ross and Fletcher 1998) and is associated with a poor prognosis and an aggressive phenotype (Slamon et al. 1987).

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By introducing trastuzumab (Herceptin) based therapy (Baselga et al. 1996) for the treatment of both, early (Slamon et al. 2011, Gianni et al. 2012) and metastatic HER2-positive breast cancer (Slamon et al. 2001), the prognosis of patients has substantially improved.

Trastuzumab works through multiple mechanisms to inhibit tumor growth, including inhibition of downstream signalling by blocking either HER2 homodimerization (Ghosh et al. 2011) or ligand-
independent HER2/HER3 heterodimerization (Junttila et al. 2009). Trastuzumab also inhibits HER2 activation by inhibiting the cleavage of its extracellular domain, thus preventing the formation of the truncated and very active form of HER2, p95HER2 (Molina et al. 2001, Nahta and Esteva 2006). Finally, trastuzumab induces antibody-dependent cell-mediated toxicity (Hudis 2007).

Despite its effectiveness in both adjuvant and metastatic settings, therapeutic resistance to trastuzumab remains an important clinical issue. Approximately, 15% of patients relapse after treatment (Kümler et al. 2014) because of acquired resistance (Pohlmann et al. 2009, Chung et al. 2013). Indeed, median time to progression remained less than one year when trastuzumab was combined with chemotherapy (Slamon et al. 2001, Esteva et al. 2002).

It is important to mention that tumor cells with acquired resistance to trastuzumab continue to depend on the HER2 oncogene. Indeed, gene amplification and RNA/protein overexpression are still present in trastuzumab-resistant HER2+ clones (Ritter et al. 2007).

The mechanisms underlying the frequent development of resistance to trastuzumab are only starting to be understood and are still under active investigation (Garrett and Arteaga 2011). Several of the proposed mechanisms of resistance to trastuzumab involve persistence or reactivation of the PI3K signalling through amplification of alternative tyrosine kinase receptor and/or mutations in the PI3K components (Rexer and Arteaga 2013). The formation of insulin-like growth factor-I receptor (IGF-1R)/HER-2 heterodimer may also contribute to trastuzumab resistance (Nahta et al. 2005). In particular, trastuzumab does not seem to be able to prevent ligand-activated HER2/HER3 or HER2/HER1 heterodimerization which could give tumor cells a way to escape from the inhibitory effects of trastuzumab (Ghosh et al. 2011).

The mechanism of HER2 signalling has been the focus of extensive research in order to identify additional targets therapies for patients with trastuzumab-resistant breast cancer. A number of agents targeting various downstream components of the pathways associated with HER2 signalling are currently under clinical investigation. These molecules include extracellular targeted therapies (monoclonal antibodies directed against HER family receptors), intracellular targeted therapies (tyrosine kinase inhibitors) and agents that target downstream effectors including members of either the mitogen-activated protein kinase (MAPK) or the phosphatidylinositol 3-kinase (PI3K) pathways. In addition to PI3K/AKT modulators, targeted therapies directed at the Hsp-90 apoptotic pathway as well as factors modulating angiogenesis are also currently being developed (Rosen et al. 2010).

THE EGFR FAMILY RECEPTORS

HER2 is one of the four members of the human EGFR family, which also includes EGFR (HER1 or ErbB-1), HER3, and HER4 (Hudis 2007). The HER gene family is encoded by genes on different chromosomes and regulate normal breast growth and development. However, their deregulations leading to the activation of downstream pathways appear to be particularly important, not only for tumor development but also for treatment efficacy (Roskoski Jr 2004).

Each HER receptor shows an extracellular domain, a helical transmembrane segment, and an intracellular protein tyrosine kinase domain. The extracellular region of each HER receptor contains four domains (I-IV). Domains I, III and IV are involved in ligand binding. The domain II loop, the so called dimerization arm, promotes direct receptor-receptor interaction.

Most of the HER receptors have a ligand. For example, HER3 has a specific ligand, heregulin (HRG), but does not have any kinase activity. HER2 is unique since it does not have any known ligand (orphan receptor) but shows a tyrosine kinase activity (Kim et al. 1998). HER2 is activated
through homo- or heterotypic interactions of its extracellular domain with that of other HER receptors (Arribas et al. 2011).

In the absence of ligand binding, HER1 and HER3 exist in a tethered conformation in which intramolecular interaction between domains II and IV blocks the function of dimerization domain II. Ligand binding to HER receptors changes the tethered conformation into an extended conformation, which exposes domain II, allowing them to undergo dimerization (Hynes and Lane 2005, Gala and Chandarlapaty 2014, Burgess et al. 2014).

Interestingly, the structure of the HER2 extracellular region dramatically diverges from those of either HER1 or HER3. The HER2 extracellular region has indeed a fixed conformation and the domain II loop is exposed to interactions (Lemmon and Schlessinger 2010). This constitutive exposure of its dimerization arm might explain why HER2 is the favored partner for the other ligand-dependent HER (Graus-Porta et al. 1997, Yarden and Sliwkowski 2001, Badache and Hynes 2004). Moreover, the high level of HER2 present at the surface of HER-positive tumor cells facilitates a spontaneous formation of HER2 homodimers (Yarden and Sliwkowski 2001, Atalay et al. 2003).

The role of HER2 in heterodimers formation seems to be related to its capacity to act as a co-receptor, increasing the affinity of the ligand binding to the heterodimerized receptor complexes (Atalay et al. 2003, Graus-Porta et al. 1997).

The HER receptors dimerization and phosphorylation lead to the activation of intracellular signalling cascades including both the phosphatidylinositol triphosphate kinase (PI3K)/protein kinase B (Akt) and the mitogen-activated protein kinase (MAPK)/ERK pathways (Yarden and Sliwkowski 2001, Atalay et al. 2003, Park et al. 2008, Rosen et al. 2010). Ultimately, these signalling cascades lead to the expression of target genes that regulate various cellular processes influencing growth, proliferation, migration and survival (Yarden and Sliwkowski 2001) (Fig. 1).

Preclinical works have reported that the HER3-ligand binding HRG enhanced HER2/HER3 heterodimer formation. Indeed, HRG induces recruitment of HER3 to an HER2-Src, resulting in upregulation of tyrosine phosphorylation and kinase activation (Vadlamudi et al. 2003, Ghosh et al. 2011). In contrast to HER3/HER2 heterodimers, HER2 homodimers and HER2/EGFR heterodimers do not induce Src kinase phosphorylation and activation (Vadlamudi et al. 2003, Huang et al. 2010).

It has been suggested that the formation of the HER2-HER3 heterodimer has the strongest transforming capacity compared to the other HER homo- and heterodimers. Despite the absence of tyrosine kinase activity, HER3 has multiple PI3K docking sites on its cytoplasmic domain, which therefore render it a potent activator of the tyrosine kinase enzyme (Xia et al. 2004).

**NOVEL STRATEGY TO TARGET HER2: PERTUZUMAB**

Pertuzumab is a recombinant, humanized, monoclonal antibody that binds to the extracellular dimerization domain II of HER2 (located on the opposite side of the domain IV where trastuzumab binds). Pertuzumab inhibits heterodimerization of HER2 with EGFR, HER3, HER4 (Nahta et al. 2004, Agus et al. 2002, Metzger-Filho et al. 2013) and IGF-1R (Nahta et al. 2005), whereas trastuzumab is preferentially active against tumors driven by HER2 homodimers (Ghosh et al. 2011).

More specifically, pertuzumab prevents ligand-induced dimerization of HER2 with HER3, thus inhibiting the activation of downstream cell signalling pathways that are critical for the tumor growth (Agus et al. 2002, Cho et al. 2003, Yarden and Sliwkowski 2001, Harari and Yarden 2000).

Pertuzumab can inhibit tumor cell growth following HRG-induced HER2 heterodimerization (Ghosh et al. 2011). However, pertuzumab, in contrast to trastuzumab, is not capable of preventing the formation of the p95HER2 truncated form (Molina et al. 2001).
For the above mentioned reasons, the different ways trastuzumab and pertuzumab work are likely to be complementary, and provide, when combined, a more complete blockade of HER2 downstream signaling than either agent alone (Fig. 2).

**PRECLINICAL DATA ON PERTUZUMAB**

*In vivo* preclinical studies showed that pertuzumab is active against various tumor types, including breast (Agus et al. 2002), ovary (Mullen et al. 2002), and others.
Suggested. When both BT474MI and MCF-7 breast cancer cell lines were treated with heregulin and either pertuzumab or trastuzumab, pertuzumab was more efficient in disrupting ligand-mediated HER2/HER3 complex formation and in blocking the appearance of a HRG-dependent phosphorylation signal of HER2 (Agus et al. 2002).

More recently, other works showed that inhibition of HER2 phosphorylation by tyrosine kinase inhibitors (TKIs) targeting EGFR and HER2 in HER2-positive breast cancer cells, lead to feedback upregulation of activated HER3, thus limiting the inhibitory effect of HER TKIs (Sergina et al. 2007, Amin et al. 2010). Indeed, RNAi knockdown of HER3 or treatment with the HER3 neutralizing antibody AMG-888, sensitized 2007), non-small cell lung carcinoma (Sakai et al. 2007), prostate (Agus et al. 2002, Mendoza et al. 2002) and colon cancer (Pohl et al. 2009).

The resolution of the 3D structure of HER2 bound to pertuzumab (Franklin et al. 2004) provided important information regarding the role played by the different domains of the receptor for its activity in the context of either its overexpression or its ligand-induced activation.

The ability of pertuzumab to inhibit in vitro tumor cell growth by blocking HER2/HER3 heterodimerization and its ligand-induced activation is a unique trait which is not shared by trastuzumab (Lee-Hoeflisch et al. 2008). The essential role of HER3 in tumor progression and drug resistance in HER2-dependent cells has previously been suggested. When both BT474MI and MCF-7 breast cancer cell lines were treated with heregulin and either pertuzumab or trastuzumab, pertuzumab was more efficient in disrupting ligand-mediated HER2/HER3 complex formation and in blocking the appearance of a HRG-dependent phosphorylation signal of HER2 (Agus et al. 2002).

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HER2-positive cells to lapatinib-induced apoptosis (Garrett et al. 2011).

In agreement, the combination of pertuzumab with trastuzumab showed a synergistic effect when compared to pertuzumab single therapy in vitro on HER2-overexpressing BT474 breast cancer cell lines (Nahta et al. 2004). The combination of the two antibodies reduced by 60% the cell survival at doses for which individual drugs did not affect it. These data were confirmed in vivo in HER2-positive breast and non-small cell lung cancer xenografts where the combination of trastuzumab and pertuzumab strongly enhanced the antitumor effect of both compounds and induced tumor regression in both xenograft models (Scheuer et al. 2009).

The enhanced efficacy of pertuzumab based combination was also observed when tumor progression occurred during the course of trastuzumab monotherapy (Scheuer et al. 2009). Here, the authors treated mice bearing KPL-4 (a human cell line isolated from the malignant pleural effusion of a breast cancer patient with an inflammatory skin metastasis; Kurebayashi et al. 1999) tumor xenografts with trastuzumab until tumor mass started to progress again (day 35). At that point, the authors treated mice with either trastuzumab alone or trastuzumab combined to pertuzumab. The combination of the two antibodies was able to inhibit tumor growth and even reduce tumor mass for an additional 45 days.

Finally, it has been shown that both trastuzumab and pertuzumab induce the activation of the antibody-dependent cellular toxicity (ADCC) pathway, which is part of their antitumor activity (Scheuer et al. 2009, Mamidi et al. 2013). Indeed, through an in vitro ADCC assay, Scheuer and colleagues showed that both trastuzumab and pertuzumab applied as a single agent effectively activated ADCC with equal potency. However, there was no increase in ADCC efficiency when both agents were combined.

**CLINICAL STUDIES**

A single-arm phase II study evaluated pertuzumab in patients who had received up to 3 trastuzumab-containing regimens and found a 24% positive response rate with 50% of patients demonstrating stable disease. This study suggested a role of pertuzumab in treating trastuzumab-resistant HER2 breast cancer (Baselga et al. 2010).

Although pertuzumab alone seemed to have antitumor activity, the combination of pertuzumab with trastuzumab was shown to be more efficient than pertuzumab monotherapy (Cortés et al. 2012). In this trial, pertuzumab was given as monotherapy to patients with advanced HER2 positive breast cancer whose disease had progressed during prior trastuzumab-based therapy. When progressive disease or unacceptable toxicity was observed, trastuzumab was reintroduced and patients received a combination of pertuzumab and trastuzumab. Progression-free survival was increased in the combination arm compared to the pertuzumab monotherapy arm (17.4 v 7.1 weeks, respectively). Importantly, the treatment was well tolerated with minimal cardiac dysfunction.

The efficacy of adding pertuzumab to trastuzumab plus docetaxel for the first-line treatment of HER2-positive metastatic breast cancer was demonstrated in a randomized, double-blind, multinational, phase III CLEOPATRA (Clinical Evaluation of Pertuzumab and Trastuzumab) trial (Baselga et al. 2012). Patients with metastatic HER2-positive breast cancer were randomly assigned to receive either pertuzumab by intravenous infusion (840 mg initial dose, 420 mg every 3 weeks thereafter) or matched placebo as an add-on to the standard-of-care trastuzumab (8mg/Kg initial i.v. dose, 6mg/Kg i.v. every 3 weeks thereafter) and docetaxel (75mg/m² i.v. every 3 weeks for at least 6 cycles, with the option of dose escalation to 100 mg/m²) (Fig. 3). A prolongation of progression-free survival (PFS) in the pertuzumab arm (18.5 vs
12.4 months, p < .001) with an objective response rate of 69.3% in the control arm compared with 80.2% in the pertuzumab arm was observed. Follow-up data at a median of 50 months showed a significant improvement of the overall survival (OS) with pertuzumab, trastuzumab, and docetaxel in patients with HER2-positive metastatic breast cancer, compared to patients receiving placebo, trastuzumab, and docetaxel (Swain et al. 2013).

On the basis of this clinical trial, pertuzumab has been approved by the U. S. Food and Drug Administration for use in combination with trastuzumab and docetaxel for the treatment of patients with HER2-positive metastatic breast cancer who have not received prior anti-HER2 therapy or chemotherapy for metastatic disease.

The positive results of the CLEOPATRA clinical trial are even more encouraging with respect to ongoing investigations on pertuzumab and trastuzumab in combination with a taxane. One example is the PERUSE trial (ClinicalTrials.gov Identifier NCT01572038), evaluating pertuzumab and trastuzumab in combination with paclitaxel, docetaxel, or nab-paclitaxel (Abraxane) with the aim to clarify whether one taxane is more appropriate than the other in the first line setting for HER2 positive advanced breast cancer.

In the first line setting, however, the combination of pertuzumab, trastuzumab with ado-trastuzumab emtansine (an antibody-drug conjugate consisting of the monoclonal antibody trastuzumab linked to the cytotoxic agent DM1) may be preferable. In that regard, the ongoing trial MARIANNE (ClinicalTrials.gov Identifier NCT01120184) has the potential to change first-line therapy for HER2 positive metastatic breast cancer. Even if both treatments are equally efficacious, the toxicity with ado-trastuzumab is minimal. T-DM1 is a novel antibody-drug conjugate incorporating the trastuzumab with the cytotoxic activity of the microtubule-inhibitory agent DM1 (a derivative of maytansine). T-DM1 has been shown to significantly prolong progression-free and overall survival with less toxicity than lapatinib plus capecitabine in patients with HER2-positive advanced breast cancer previously treated with trastuzumab and a taxane (Verma et al. 2012).

Based on exciting results of trials which evaluated pertuzumab in the neoadjuvant setting, the drug was granted FDA-accelerated approval in 2013. In this setting, pathological response rates of the combination of pertuzumab or trastuzumab, or both, with docetaxel and the combination of pertuzumab and trastuzumab without chemotherapy were evaluated in the NeoSphere trial. Patients with locally advanced inflammatory or early breast cancer were treated and a significant improvement in pathological complete response (pCR) was
observed in the group of patients given pertuzumab and trastuzumab plus docetaxel compared with those given trastuzumab plus docetaxel (Gianni et al. 2012). Interestingly, in patients treated with pertuzumab, trastuzumab plus docetaxel, the pCR rate was higher in patients with estrogen receptor (ER)-negative tumors (63.2%) compared with those with ER+ tumors (27.3%) (Gianni et al. 2012).

In the TRYPHAENA (Trastuzumab Plus Pertuzumab in Neoadjuvant HER2-Positive Breast Cancer) trial, patients with HER2+ tumors were randomized in three arms to receive six neoadjuvant cycles q3w (Arm A: 5-fluorouracil, epirubicin, cyclophosphamide [FEC] followed by docetaxel [T]; with trastuzumab [H] and pertuzumab [P] given concurrently throughout [FEC + H + P x 3 → T + H + P x 3]; Arm B: FEC followed by T + H + P [FEC x 3 → T + H + P x 3]; or Arm C: T, carboplatin; H with P [TCH + P x 6]. Following neoadjuvant therapy, patients underwent surgery and continued trastuzumab to complete 1 year of treatment (Fig. 4). The primary end-point was cardiac safety. All grades of symptomatic left ventricular systolic dysfunction (LVSD) were low across all 3 study arms: 5.6%, 4.0%, and 2.6% in Arms A, B, and C, respectively. Grade 3 or higher LVSD was observed in 2.7% of patients in Arm A but not in Arms B and C. In conclusion, the combination of pertuzumab with trastuzumab and the standard chemotherapy resulted in low rates of symptomatic LVSD (Schneeweiss et al. 2013).

The combination of pertuzumab and trastuzumab is also under investigation in the adjuvant setting. The APHINITY trial (Adjuvant Pertuzumab and Herceptin in Initial Therapy of breast cancer; ClinicalTrials.gov Identifier NCT01358877) is a placebo-controlled study in patients with HER2-positive primary breast cancer who have had an excision of their tumor. Patients were randomized in two arms: (1) the investigational arm with a course of adjuvant chemotherapy consisting of a taxane-based regime (anthracycline-taxane or taxane-platin) and trastuzumab and pertuzumab for 1 year; (2) the comparator arm which consisted in the same adjuvant chemotherapy with trastuzumab and placebo for 1 year. The primary objective was to compare invasive disease-free survival between both treatment arms.

| Neoadjuvant regimen | Adjuvant regimen |
|---------------------|------------------|
| Cycles 1-3          | Cycles 4-6       |
| Arm A               |                  |
| H + Pertuzumab      | H                |
| FEC                 | T                |
| Arm B               |                  |
| FEC                 | T + H + Pertuzumab |
| Arm C               |                  |
| TCH + Pertuzumab    | H                |

**Figure 4 - The TRYPHAENA study design.** The trial evaluated trastuzumab plus pertuzumab in neoadjuvant HER2-positive breast cancer. H: Trastuzumab; FEC: 5-fluorouracil, epirubicin, cyclophosphamide; T: docetaxel; TCH = docetaxel, carboplatin, and trastuzumab.
CONCLUSIONS

The introduction of trastuzumab in the treatment of HER2-positive metastatic breast cancer patients favorably changed the natural history of this disease. However, despite this major advance, HER2-positive metastatic breast cancer will eventually progress in most patients. One of the mechanisms responsible for the development of resistance to trastuzumab is the heterodimerization of HER2 with other HER receptors which may redundantly trigger cell proliferation signals.

A few years ago, one might have asked why we should combine two therapeutic antibodies targeting the same receptor. Today, in view of the mechanistic differences that have been identified through numerous preclinical works between trastuzumab and pertuzumab, this therapeutic approach should be encouraged. Pertuzumab, which could be considered as a prototype of HER2 dimerization inhibitor, shows mechanistic advantages that distinguish it from trastuzumab, in particular in regards to HER2 heterodimerization. However, one might consider that trastuzumab also shows mechanistic advantages over pertuzumab in regards to its ability to prevent the formation of the p95HER2 truncated form of HER2, thus highlighting their functional complementarity. This particularity is likely to play an important role on the demonstrated synergistic effect when both compounds are combined.

The approval of pertuzumab marked the first licensed dual anti-HER2 regimen for treatment of breast cancer and is likely to represent a major advance in the treatment of this pathology, comparable to the approval in 1998 of trastuzumab. The clinical benefit of dual treatment with HER2-targeted antibodies with complementary mechanism of growth inhibition should also encourage further research in this field and open new therapeutic strategies. Consequently, numerous molecules are currently being developed. For example, LJM716 is a novel monoclonal antibody that is capable of neutralizing either ligand-dependent or independent HER3 signalling by locking HER3 in its inactive conformation. LJM716 is a potent inhibitor of both HER3/AKT phosphorylation and proliferation in HER2-expressing cancer cells and has displayed single agent efficacy in tumor xenograft models (Garner et al. 2013).

The development of a bispecific anti-HER2 antibody using both trastuzumab and pertuzumab is another example. This bispecific antibody (named TPL) retained the full binding activities of both parental antibodies, and exhibited pharmacokinetic properties similar to those of a conventional IgG molecule. TPL showed superior HER2 heterodimerization-blocking activity over the combination of both parental monoclonal antibodies. The unique potential of TPL to overcome trastuzumab resistance should be considered as a promising treatment in the clinic (Li et al. 2013).

Despite these recent advances, there are still a number of issues to be addressed. In particular, the identification of biomarkers is needed to identify patients more likely to respond, and to avoid treating patients likely to experience a worse outcome compared to the standard of care.

To date, and despite multiple researches in this area, the only robust biomarker allowing the prediction of response to HER2-targeted therapies is HER2 itself. Currently, HER2 status is determined by measurement of HER2 receptor protein and/or erbB2 gene amplification by immunohistochemistry or fluorescence in situ hybridization (FISH) (Singer et al. 2008). While the negative predictive value of these assays for predicting the absence of benefit from trastuzumab-based therapy is high, their positive predictive value remains insufficient. The clinical benefit and response rates appear to depend on the intensity of HER2 overexpression (2+ or 3+) with response rates of 35% in grade 3+ expressors compared to only minimal benefit in 2+ positives (Vögel et al. 2002).
Results of biomarker analysis in the CLEOPATRA (Baselga et al. 2014) trial were consistent with those with smaller TRYPHAENA and NeoSphere of neoadjuvant trastuzumab plus pertuzumab in patients with HER2 positive early breast cancer, where the primary efficacy endpoint was pCR (Schneeweiss et al. 2012, Gianni et al. 2011). The PI3KCA status identified a subpopulation of patients with HER2-positive disease with poor prognosis when treated with anti-HER2 antibodies. Based on the findings of CLEOPATRA trial, clinical trials of HER2-targeted molecules in combination with PI3K pathway targeted agents may therefore be justified.

The identification of biomarkers is thus, urgently required in order to enable a better care of patients. Also, the medical staff would be able to adapt the treatment on the basis of the molecular features of the tumors and anticipate their modifications during the course of tumor progression. The understanding of the underlying mechanisms of either intrinsic or acquired resistance to HER2-targeting antibodies is also necessary, to define new therapeutic strategies and identify new targets. The ongoing development and success of these novel approaches rely, more than ever, on translational collaborations between basic scientists, preclinical researchers and clinicians.

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