



Is it useful to combine taxanes with targeted agents in mCRPC patients – have we hit the target in this phase II study with docetaxel and curcumin allowing going for a phase III study?

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Ten percent to 20% of men diagnosed with prostate cancer will eventually develop castration-resistant prostate cancer (CRPC). Standard treatment for symptomatic metastatic CRPC (mCRPC) has consisted of chemotherapy-based regimens in combination with steroids for the past two decades. Docetaxel with daily prednisone is the recommended first-line therapy for symptomatic mCRPC (1). Docetaxel is the first treatment to have shown an increase in median overall survival (OS) for mCRPC. With a gain of 2–3 months OS (2–4), three key phase II randomized or III studies demonstrated the clinical benefit of docetaxel over the previous standard treatment, the mitoxantrone, an anthracedione (1) (*Table 1*).

However, only half of the patients will have an objective response with a median OS of 2 years before the emergence of new hormone therapy treatments. Recently, highly effective novel therapies have been approved. Recent clinical trials focus on treatment before or after docetaxel. Five new therapies demonstrated an improvement in OS (5), three of them pre-docetaxel administration: sipuleucel-T (6), enzalutamide (7), abiraterone acetate (8) and four treatments in the post-docetaxel setting: radium-223 dichloride (9), abiraterone acetate (10), enzalutamide (11) and cabazitaxel (12). Cabazitaxel is the only drug to have been compared to docetaxel in a phase III trial; the results of the FIRSTANA study showed similar OS with both taxanes but different toxicity profiles (13). Combinations of docetaxel with new targeted agents have been studied in order to increase its therapeutic effects but without success, mainly because

phase I went directly to phase III without selecting a specific population to increase the efficacy of the combination therapy of docetaxel plus targeted agent (*Table 2*).

Mahammedi *et al.* reported the results of a phase II pilot study assessing the combination docetaxel, prednisone and curcumin (23). Pre-clinical studies suggested that curcumin increased docetaxel/prednisone cytotoxicity through downregulation of various cell cycle regulatory proteins (24), and seemed to provoke cell cycle growth arrest at the G1/S phase, by downregulation of cyclins (25). Sun *et al.* also suggested that the association of curcumin and docetaxel had pharmacokinetics propriety and increased taxanes therapeutic efficacy via the inhibitory role of curcumin on hepatic organic anion transporting polypeptide (OATP) 1B1, OATP1B3 and microsome activities (26). Thirty patients with mCRPC received docetaxel 75 mg/m² every 3 weeks with dexamethasone premedication, prednisone 5 mg BID and 6,000 mg of curcumin daily for 7 consecutive days (from day –4 to day +2 of each cycle), as determined in the preliminary phase I study (27). An objective PSA response was observed in 59% (n=17) of patients. The objective response rate (ORR) was 40% (n=6) among the 15 patients with evaluable lesions. Those results were similar to those observed in the three main studies on docetaxel (2–4). There is no evidence to believe that this combination is more effective than docetaxel-prednisone alone in symptomatic mCRPC patients. Thus, those results are insufficient to recommend a phase III study.

An ancillary study assessed chromogranin A (CgA) and neuron-specific enolase (NSE) seric values, attempted to

Table 1 Key studies demonstrating improved overall survival with docetaxel

Trial design	Treatment	N	OS (95% CI) (months)	PSA decrease >50%	Median PSA TTP (95% CI) (months)	References
Phase III	A: docetaxel 75 mg/m ² + prednisone 5 mg bid, D1=D21; B: docetaxel 30 mg/m ² weekly for 5/6 weeks + prednisone bid; C: mitoxantrone 12 mg/m ² + prednisone 5 mg bid, D1=D21	A=335; B=334; C=337; total: 1,006	A: 18.9 (17.0–21.2); B: 17.4 (15.7–19.0); C: 16.5 (14.4–18.6)	A: 45%; B: 48%; C: 32%; P<0.001	A: 7.7 (7.1–8.6); B: 8.2 (6.3–11.5); C: 7.8 (5.4–10.5)	Tannock <i>et al.</i> [2004] (2)
Phase III	A: docetaxel 60 mg/m ² on day 2 + estramustine 280 mg tid on days 1–5 + 60 mg of dexamethasone before docetaxel, D1=D21; B: mitoxantrone 12 mg/m ² + prednisone 5 mg bid, D1=D21	A=338; B=336; total: 674	A: 17.5; B: 15.6; P=0.02	A: 50% (n=155); B: 27% (n=82)	A: 6.3; B: 3.2; P<0.001	Petrylak <i>et al.</i> [2004] (3)
Phase II randomized	A: docetaxel 70 mg/m ² on day 2 + estramustine (280 mg PO tid on days 1–5 and 8–12) + prednisone 10 mg daily, D1=D21; B: docetaxel 35 mg/m ² on days 2 and 9 + estramustine (280 mg PO tid on days 1–5 and 8–12) + prednisone 10 mg daily, D1=D21; C: mitoxantrone 12 mg/m ² + prednisone (10 mg daily), D1=D21	A=43; B=42; C=42; total: 130	A: 18.6 (14.9–22.3); B: 18.4 (14.1–22.8); C: 13.4 (9.4–17.5)	A: 67% (n=29); B: 63% (n=26); C: 18% (n=7); P<0.0001	A: 8.8 (6.9–10.8); B: 9.3 (7.5–11.1); C: 1.7 (0.7–2.7); P=0.000001	Oudard <i>et al.</i> [2005] (4)

N, number of patients; OS, overall survival; TTP, time to progression; 95% CI, 95% confidence interval.

Table 2 Phase III combination therapies with docetaxel

Agent studied	Number of patients	Docetaxel OS (months)	Docetaxel + agent OS (months)	References
Calcitriol	954	20.2	17.8	Scher <i>et al.</i> [2011] (14)
GVAX vaccine	408	14.1	12.2	Higano <i>et al.</i> [2009] (15)
Atrasentan	994	17.6	17.8	Quinn <i>et al.</i> [2013] (16)
Zibotentan	594	19.2	20.0	Fizazi <i>et al.</i> [2013] (17)
Bevacizumab	1,050	21.5	22.6	Kelly <i>et al.</i> [2012] (18)
Lenalidomide	1,059	Not reached	17.7	Petrylak <i>et al.</i> [2015] (19)
Dasatinib	1,522	21.2	21.5	Araujo <i>et al.</i> [2013] (20)
Afliberceptp	1,224	21.2	22.1	Tannock <i>et al.</i> [2013] (21)
Custirsen	1,022	22.2	23.4	Chi <i>et al.</i> [2015] (22)

OS, overall survival.

define prognostic markers to treatment response. CgA and NSE are frequently used as markers of neuro-endocrine (NE) differentiation. NE differentiation is correlated with a more aggressive disease (28). However, the prognostic role of CgA and NSE is unclear and its use in therapeutic decisions has not yet been defined. Twelve out of 22 patients had elevated baseline CgA levels, 6 NSE and 4 both markers. Those markers were correlated with PSA rate before treatment: patients with lower levels of PSA had significantly higher levels of NSE and lower CgA values. ORR and PSA response for patients with elevated CgA and NSE were the same as patients with no NE markers. However, NE variation was different according to the markers: a higher decrease rate was observed for NSE. The treatment seemed to be more active on patients with NSE marker than CgA. CgA and PSA had parallel evolutions. Before considering any phase III study for the combination of docetaxel, prednisone and curcumin, prognostic biomarkers are needed to define an eventual subgroup of mCRPC patients who could benefit from it.

Optimization of the docetaxel-prednisone combination is needed. However, past experiences of negative phase III with combination therapies are numerous (29). Finding a unique combination therapy effective for all patients is probably not realistic. Rather than associating new molecules with docetaxel in order to improve its response rates research might focus on understanding the biological differences in patients with mCRPC, enlightening the mechanism of response to docetaxel and identifying a targeted population who would benefit from this treatment. As it has been done with sunitinib and clear cell renal cell metastatic

carcinoma (30), a translational clinical study identifying subgroups of patients based on proteomic and genomic analyses would be beneficial for a more optimal treatment. To our knowledge, translational clinical studies have yet to be lead in docetaxel treatment based regimens for mCRPC.

While resistance to androgen therapies via the detection of androgen receptor (AR) splice variant 7 messenger RNA (ARV7) in circulating tumor cells (CTC) has been investigated, predictive biomarkers of response or resistance to docetaxel in mCRPC are lacking (31). Recently, researchers have been able to understand various mechanism of resistance to docetaxel (32), such as an activated AR, activated tyrosine kinase receptors (RTK), anti-apoptotic signaling, aberrant proangiogenesis, unfavorable microenvironment, increased drug efflux, mutation of the drug target or drug resistant cancerous cell populations as NE differentiated cells.

A single drug combination will therefore be insufficient to encounter the diversity of these mechanisms. Multiple targets for new therapies are now being tested (29). Selecting the proper population should be a major concern in clinical trials. As molecular biology is making progress, more and more mechanism of resistance are identifying. In this light, clusterin is a chaperone molecule whose inhibition increased cell-sensitivity to many cytotoxic agents, including taxanes, by delaying the end of the mitotic phase. However, the Clusterin inhibitor custirsen failed to improve OS when combined with docetaxel. Further molecular compensatory mechanisms have recently been identified: the upregulation of cell cycle involved kinase Wee1. Simultaneous inhibition of Clusterin and Wee1 may improve synergistic responses

of combinatorial regimens using taxanes (33).

Moreover, recent data suggested partial cross-resistance between taxanes and AR targeting agents (abiraterone acetate and enzalutamide) (34). On the other hand, some patients might benefit from a re-challenge with docetaxel. These results raise the question of treatment sequencing with three effective therapies before docetaxel and four approved treatments after docetaxel administration, determining the best sequence is still an issue.

In order to improve mCRPC survival, we should concentrate on understanding the diversity of the biological mechanism involved in the tumour-cell responses to treatment and on optimizing the therapeutic sequences in order to personalize the standard of care to patients' specific cancer biology.

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