

Material and Methods: From July 2010 to April 2011 the NIBIT-M1 enrolled the 86 cutaneous MM pts, stage III (2) or IV (84) pts (60 males, 26 females), median age 54 (24–78) years, ECOG performance status 0–1, planned in the study. Twenty-one pts had evidence (19) or history (2) of brain metastases. Forty-three pts were treatment naive and the remaining had received one line of systemic treatment for metastatic disease. Ipilimumab was administered i.v. at 10 mg/kg q3 weeks (wk) for 4 doses in the induction phase (IP) and once q12 wk from wk 24 in the maintenance phase (MP); fotemustine was administered i.v. at 100 mg/m² weekly for 3 wk (IP), and q3 wk from wk 9 (MP). Tumour assessment (TA) *per* immune-related response criteria (irRC), was performed at screening and wk 12, then every 8 wk until W36, and every 12 wk from W36 onwards. A pre-spe-specified safety analysis was planned at wk 6 of treatment for the initial 18 pts. Adverse Events (AE) and immune related AE (irAE) were collected according to Common Terminology Criteria for Adverse Events version 4.0.

Results: On November 2010, the safety analysis was successfully met, and no additive toxicities were observed; thus, the Safety Committee allowed resuming the accrual. As of April 2011, 28 pts have terminated the IP and 13 of them have already entered the MP. Of the remaining pts, 43 are completing the IP and 15 have been withdrawn for AE severity (1) or disease progression (14). Of the 17 pts for which TA at wk 12 is available, 14 achieved a disease control (CR, PR or SD), including brain metastases in 5 out of 6 pts, while 3 had PD.

Conclusions: Though very initial, the available data suggest for the safety and efficacy of ipilimumab in combination with fotemustine in MM pts with or w/o brain metastases. The six months results from the study closure will be presented.

9306

ORAL

Safety and Efficacy of Ipilimumab-treated Patients With Melanoma and Brain Metastases

K. Margolin¹, F.S. Hodi², D.F. McDermott³, D.P. Lawrence⁴, O. Hamid⁵, I. Puzanov⁶, J.A. Thompson⁷, M.S. Ernstoff⁸, T.L. Michener⁹, K.N. Heller¹⁰. ¹University of Washington, Division of Medical Oncology, Seattle, ²Dana-Farber Cancer Institute, Department of Medicine, Boston, ³Beth Israel Deaconess Medical Center, Department of Cell Biology, Boston, ⁴Massachusetts General Hospital Cancer Center, Department of Hematology/Oncology, Boston, ⁵The Angeles Clinic and Research Institute, Neuro-Oncology Clinic, Santa Monica, ⁶Vanderbilt University Medical Center, Department of Oncology, Nashville, ⁷University of Washington, SCCA Melanoma Clinic, Seattle, ⁸Dartmouth-Hitchcock Medical Center, Department of Hematology/Oncology, Lebanon, ⁹Bristol-Myers Squibb Company, Collaborative Science Center of Excellence, Plainsboro, ¹⁰Bristol-Myers Squibb Company, Oncology Medical Strategy, Plainsboro, USA

Background: Ipilimumab (YervoyTM), an anti-CTLA-4 monoclonal antibody that augments T-cell-mediated antitumour responses, is indicated in the US for patients (pts) with unresectable or metastatic melanoma. At diagnosis of metastatic disease, 30% of pts have brain metastasis and an additional 30% will develop these within 1–2 years. Herein, we describe overall survival and safety of ipilimumab in pts with melanoma and brain metastases in clinical trials CA184–042 (NCT00623766) and CA184–045 (NCT00495066).

Patients and Methods: The prospective trial CA184–042 included pts with active, measurable brain metastasis (≥1 lesion >0.5 cm and/or ≥2 lesions >0.3 cm with none >3 cm). At baseline, pts were either stable without steroid therapy (Arm A) or required steroids for central nervous system symptoms (Arm B). Ipilimumab 10 mg/kg was given Q3W for four doses with potential maintenance dosing Q12W. The expanded access program CA184–045, a multicenter, open-label study of ipilimumab 3 or 10 mg/kg Q3W for four doses, included pts with stable and asymptomatic brain metastases at baseline. Among pts who received 10 mg/kg, overall survival (OS) at 1 year was retrospectively collected via database; pts lost to follow up were assumed dead. Safety was monitored prospectively in both trials.

Results: In CA184–042, 51 pts in Arm A and 21 in Arm B were treated with ipilimumab 10 mg/kg. Patients in Arms A & B were all Caucasian with ECOG-PS of 0 or 1, 65% and 52% male, and of mean age 58 and 55, respectively. The 12- and 18-month OS in pts not requiring steroids was 30% at both time points (CIs 0.2–0.5; 0.2–0.4, respectively). The 12-month OS rate in pts with symptomatic brain metastases was 10% (CI 0.0–0.3). There were no unexpected toxicities – grade 3–4 central nervous system adverse events (AEs) occurred in 31% of patients in Arm A and 29% in Arm B. In CA184–045, of 874 pts treated with ipilimumab 10 mg/kg, 165 were identified with brain metastasis. The 1 year OS for these pts was 20%. Drug-related AEs of any grade and grade 3/4 occurred in 41% and 22% of all pts, respectively.

Conclusions: Safety and efficacy of ipilimumab in pts with melanoma and brain metastases are consistent between the prospective and open label trials and ipilimumab also shows similar antitumour activity in the brain as reported overall for extracranial metastases. Two-year survival results and safety observations from fully mature CA184–042 data will be presented.

Poster Presentations (Sun, 25 Sep, 14:00–16:30)

Melanoma and Skin Cancer

9307

POSTER

Oncogenic Mutation Dependent Response to Growth Factors in Melanoma Cells

T. Garay¹, E. Juhasz¹, J. Dobos², V. Laszlo³, W. Berger⁴, M. Grusch⁴, J. Timar¹, B. Hegedus¹. ¹Semmelweis University, 2nd Department of Pathology, Budapest, ²National Institute of Oncology, Pathology, Budapest, Hungary; ³Medical University of Vienna, Thoracic Surgery, Vienna, ⁴Medical University of Vienna, Internal Medicine I, Vienna, Austria

Background: Malignant melanoma has one of the worst prognosis among solid tumours. The high mortality of melanoma is due to the metastatic potential of tumour cells that requires increased cell motility. Epidermal and basic fibroblast growth factors (EGF and FGF2) are major autocrine and paracrine signaling molecules in human melanoma. Since two of the most common oncogenic mutations, namely BRAF and NRAS are critical components of this signaling network, in this study we compared the oncogenic mutation dependent effect of growth factors on the proliferation and migration of melanoma cells.

Materials and Methods: Growth factor receptor expression had been measured by Western blot and NRAS and BRAF mutations had been determined by direct sequencing and restriction fragment length polymorphism, respectively. Cell motility and proliferation were determined by the analysis of three-days-long time-lapse videomicroscopic recordings. Both the baseline and induced activation of the growth factor receptor pathway had been quantified by the immunoblot analysis of the phosphorylation of two major downstream effectors, including Erk1/2 and S6.

Results: Both BRAF and NRAS mutations resulted in a higher baseline activation of Erk1/2 and S6 when compared to double wild-type cells under control conditions. Both mutations attenuated the activation of the two downstream targets in response to EGF and FGF2 treatment. Interestingly we found a more profound response in cell motility as compared to cell proliferation. Of note, double wild-type cells responded to both EGF and FGF2 treatment. In contrast BRAF and NRAS mutated melanoma cells displayed varying degree of sensitivity to these growth factors.

Conclusions: In summary our findings demonstrate that the different oncogenic mutations in melanoma cells have an impact on the mitogenic effect of the activation of growth factor receptor signaling networks. Since a large number of the emerging molecularly targeted therapies aim at the growth factor receptor signaling, the appropriate mutational analysis of melanoma cases are essential in both preclinical studies and in the clinical trials and practice.

9308

POSTER

The XPC A2920C, the XPF T30028C and the P53 Arg72Pro Polymorphisms, Involved in DNA Repair, Alter the Risk for the Malignant Melanoma

C. Oliveira¹, G.J. Lourenco¹, J.A. Rinck-Júnior¹, A.M. Moraes¹, C.S.P. Lima¹. ¹State University of Campinas Faculty of Medical Sciences, Department of Internal Medicine, Campinas SP, Brazil

Background: The XPC, the XPF and the P53 genes act on the nucleotide excision repair of the UV induced DNA damage and seem to be involved in origin of malignant melanoma (MM). The variant C allele of the XPC A2920C and the wild Arg allele of the P53 Arg72Pro polymorphisms encode proteins with lower activities in DNA repair than those coded by others alleles. To the best of our knowledge, there are no functional studies of the proteins encoded by the wild and variant alleles of the XPF gene. The roles of these polymorphisms for the risk of MM are unclear and therefore this was the aim of the present study.

Material and Methods: Genomic DNA from peripheral blood of 137 consecutive MM patients and 137 age and race-matched controls were analyzed by the PCR-RFLP.

Results: The frequency of the XPC CC variant genotype (13.9% vs 6.6%, $P = 0.03$) and the P53 Arg/Arg wild-type genotype (60.6% vs 47.5%, $P = 0.02$) were higher in patients than in controls. Carriers of the genotypes had a 2.47 (95% CI: 1.05–5.84) and a 1.73 (95% CI: 1.05–2.84) fold increased risks for disease than others, respectively. The frequency of the XPC CC and P53 Arg/Arg combined genotype was higher in patients than