

Poster Session #2

Classification	#	Last Name	First Name	Title of Abstract
B	77	Slipicevic	Ana	New insights into regulation mechanisms of FABP7 and its relevance for clinical outcome in malignant melanoma
B	78	Shin	Seung-Shick	Oncogenic activities of GRM1 in melanocyte transformation
B	79	Liu	Feng	Response of microphthalmic transcription factor (MiTF) to UV and ROS stress and a possible mechanism for melanoma carcinogenesis
B	80	Arndt	Stephanie	Loss of POEM promotes tumor progression in malignant melanoma
B	81	Baillet	Olivier	Syk functions as a tumor suppressor in melanoma cells by restoring senescence-like growth arrest
B	82	Wolter	Keith S.	Autophagy as a p53-independent control of melanoma cell survival
B	83	Weeraratna	Ashani	Wnt5a regulates melanoma antigen recognized by T-cells 1 (MART-1), via changes in STAT3 phosphorylation
B	84	Besch	Robert	RIG-1 and MDA-5 activate apoptosis in melanoma via Puma and Noxa
B	85	Byron	S	Evaluation of FGFR2 in melanocytes: importance of the extracellular domain....
B	86	Carreira	Suzanne	The Brn-2 transcription factor controls melanoma proliferation through repression of Mitf
B	87	Chen	Dahu	Ski deficiency inhibits human melanoma tumor growth in vivo by restoring signatures of active TGF-beta signaling
B	88	Chen	Yih-Wen	The regulation of the immunoregulatory molecule B7H3 in melanoma cell migration and invasion
B	89	Villanueva	J	Chronic B-Raf inhibition leads to drug resistance and increased invasiveness
B	90	Corre	Sebastian	Identification and implication of a new post-translational modification of the USF1 transcription factor in melanocytes
B	91	dela Serna	Ivana	The role of SWI/SNF chromatin remodeling enzymes in melanocyte and melanoma gene expression
B	92	Dumaz	Nicolas	Phosphodiesterases mediate cross-talk between the cAMP and the RAS/RAF/MEK/ERK pathways in melanoma
B	93	Ekmekcioglu	Suhendan	MDA-7/IL-24 Protein leads to melanoma cell apoptosis via differential signaling of interferon regulatory factors
B	94	Eskandarpour	Malihe	Oncogenic NRAS has a pivotal role in the malignant phenotype of human melanoma cells
B	95	Feng	Whei	Matrix metalloproteinase-1 (MMP-1) is a critical effector of BRAF kinase activation in primary human melanocytes and melanomas

B	96	Fichtman	Boris	MAPK-independent pathways in melanoma development and progression
B	97	Guadagno	Thomas	Oncogenic B-Raf V600E drives mitotic abnormalities in melanoma cells
B	98	Haferkamp	Sebastian	Mechanisms of BRAF induced cellular senescence of primary human melanocytes
B	99	Hersey	Peter	Endoplasmic stress (ER) in melanoma cells. Friend or foe?
B	100	Hoek	Keith S.	In vivo switching of human melanoma cells between proliferative and invasive states
B	101	Hohenauer	Tobias	Increased expression of Brn3a promotes survival of melanoma cells
B	102	Hu	Rong	p53 and cyclin E1-dependent effects of Skp2 on melanoma cell cycle
B	103	Kalabis	Mizuho	Identification of the Tenascin-C as a key component in melanoma stem cell niche
B	104	Keefe	Megan	Molecular characterization of melanoma-endothelial cell communications and their functional role in tumor progression
B	105	Kumar	Suresh	BRAF V600E increases transcription of HIF-1alpha in melanoma cells
B	106	Lachiewicz	Anne	Coronin 1C expression is associated with activation of the MAP-ERK kinase pathway in melanomas
B	107	Larribere	Lionel	Molecular mechanisms of senescence in human melanocytes: Differential impact of mutations in the MAPK pathway
B	108	Lee	John T	Activated Notch1 transforms primary human melanocytes
B	109	Li	Ling	Melanocyte differentiation of multi-potent neural crest-like cells from Human dermis
B	110	Lin	Jingrong	Acquired melanocytic nevi are polyclonal
B	111	Lin	Qiushi	SKI functions as a sensor and modifier of TGF-beta signaling by switching protein complexes in human melanoma cells
B	112	Lister	James	Molecular and genetic analysis of mitfa function in zebrafish melanocytes
B	113	Uffort	Deon	Regulation of iNOS expression by the MAPK pathway is mediated by NFkB in melanoma cells
B	114	Longley	B. Jack	MAGE-A, mMAGE-b and MAGE-C proteins form complexes with KAP1 and suppress p53 dependent apoptosis in MAGE positive melanomas
B	115	Sunimoto	Hidetoshi	MITF protein levels determine the proliferation status of melanoma cells
B	116	Maddodi	N	Regulation of microtubule associated protein (MAP2) in melanoma....
B	117	Martino	JJ	Characterization of GRM1 signaling pathway components in human melanoma cells
B	118	Matin	Rubeta	p63 and p73 in malignant melanoma
B	119	McKenzie	Heather	The effect of melanoma-associated p16INK4a mutants on the function of the wild-type protein

B	120	Mueller	Daniel W.	Hugl-1 expression is indirectly regulated by microRNA miR-196a in malignant melanoma
B	121	Murtas	Daniela	Oxidative stress and its relationship with p53 and survivin proteins in cutaneous malignant melanoma
B	122	Niles	Richard M	Expression and function of HIF-1 alpha in human melanoma under non-hypoxic conditions
B	123	Noonan	Frances	UV-induced melanoma in the HGF/SF transgenic model is enhanced by deficiency in the nucleoside excision repair components XPA, XPC or XPD but not CSB
B	124	O'Connell	Michael	ROR2 modulates WNT5a mediated motility in metastatic melanoma
B	125	Palmieri	Giuseppe	Fanconi genes may contribute to melanoma pathogenesis
B	126	Pawelek	John	A new phenotype in malignant melanoma: Vesiculated Golgi complex, beta 1,6-branched N-glycosylation and autophagy
B	127	Sturm	Richard	Osteonectin downregulates E-cadherin, induces Osteopontin and focal adhesion kinase activity stimulating an invasive melanoma phenotype
B	128	Piras	Franca	Apoptosis-related factors and immune system cells in cutaneous malignant melanoma
B	129	Pontecorvi	Marco	Tbx2 recruitment of HP1-alpha to the p21 initiator: a mechanism to bypass p53 mediated senescence pathways in melanoma
B	130	Rodriguez	Mercedes	Role of T-box factors during melanoma progression
B	131	Rothhammer	Tanja	Regulation of BMP expression in malignant melanoma
B	132	Scanga	Lori	Consumption of the epidermis in melanoma is associated with basement membrane disruption assessed by collagen IV and laminin immunohistochemistry
B	133	Schepsky	Alexander	Deciphering the Mitf code: BRAF regulates Mitf acetylation to determine target gene specificity
B	134	Sinnberg	T	A new kinase in melanoma, closely connected to beta-catenin signaling and apoptosis induction
B	135	Schlegel	Natalie	ID2 is involved in TGF-beta's effects on melanoma growth and proliferation
B	136	Segura	Miguel	Role of microRNA in melanoma genesis and progression
B	137	Selfridge	Jim	Production of an improved mouse model for UV-induced melanoma
P	138	Silverthorn	Courtney	Melanoma-specific gene expression is altered by a hypoxic microenvironment: implications for therapy and radiation sensitivity
P	139	Ryu	Byungwoo	Comprehensive expression profiling of tumor cell lines identifies molecular signatures of malignant melanoma progression

P	140	Baehner	Rick	RNA yields and RT-PCR gene expression profiles obtained from manual-microdissected, fixed-paraffin-embedded tissue from malignant melanomas of varying thicknesses
P	141	Lanfrancone	Luisa	RaLP, a new member of the Shc family, and L1-CAM affect melanoma migration
P	142	Tveito	Siri	New methods for the detection and molecular characterization of micrometastatic cutaneous and uveal melanoma cells in clinical samples
P	143	Tuthill	RJ	Failure of senescence as a pathway to melanoma in the dysplasia-melanoma sequence: demonstration using a tissue microarray and immunostains
P	145	Soikkeli	Johanna	Systematic search for the best gene expression markers for the detection of melanoma micrometastases
P	146	Tsai	James	Characterizing Therapeutic Response to a Selective B-Raf ^{V600E} Inhibitor by Gene Expression Analysis in Melanoma Cell Lines
P	147	Enge	Martin	Microarray analysis to compare tissue models mimicking vertically growing melanomas
P	148	Galibert	Marie-Dominique	Melanoma Genomic-Screening
P	149	Gast	Andreas	Chromosome copy-number analysis of melanoma cell lines using high resolution SNP microarrays
P	150	Jonsson	Goran	Analysis of genomic patterns in metastatic melanoma
P	151	Ju	Jingfang	Global comparative gene expression analysis of melanoma patient samples, derived cell lines and corresponding tumor xenografts
P	152	Krauthammer	Michael	Synergistic apoptotic pathways induced by 5-Aza-2'-Deoxy-Cytidine in melanoma cells as revealed by genome-wide expression profiling
P	153	Mariotti	Agnese	Proteomics analysis of membrane rafts of melanoma cells with different malignant properties
P	154	Martins	Waleska	Alterations in cell-cell and cell-extracellular matrix interactions in human melanoma as revealed by microarray analysis
P	155	Muto	Nair Hideko	Differential expression of genes related to cell adhesion and intercellular junction and their role in the malignant transformation of nevi
P	155	Spinnler	Clemens	Microarray analysis to elucidate integrin-dependent wtp53 inactivation in malignant melanomas
P	156	Neto	Joao	Gene expression profiling of cutaneous melanoma related to clinical outcome
P	157	Ouhitit	Alal	Identification of p16-downstream target genes implicated in the early dysfunction of UV-induced cell cycle mechanisms and initiation of melanoma
P	158	Pelizzola	Mattia	Chromosome-wide methylation analysis in normal melanocytes and melanoma cells gives new insights on the effectiveness of DNA promoter methylation in transcriptional repression

P	159	Prasmickaite	Lina	Characterization of melanoma cells from lymph node metastases: relevance to tumorigenecity and metastasis
P	160	Rambow	Florian	Transcriptomic analysis of spontaneous melanoma regression in MeLiM pigs