

**Additional File 4:** Lists of the genes represented by redundant clones in the RGP and Met libraries. The tables contain the identifier, Gene Ontology annotation and available information about role in cancer related processes for the genes represented by redundant clones in the RGP and Met libraries.

**Table S3: Genes represented by a redundant number of sequences in the RGP SSH library**

Symbol	Accession number	Redundancy	GO function	Reported in cancer	Reported in melanoma
ACAT2	NM_005891	2	acyltransferase activity	upregulation in hepatocarcinoma [1]	ND
ACTB	NM_001101	3	structural constituent of cytoskeleton	actin dynamics alteration is observed in tumor progression [2]	
ACTG1	NM_001614	2	structural constituent of cytoskeleton	actin dynamics alteration is observed in tumor progression [2]	reduction of gamma-actin in invasive melanoma cells have been reported [3]
ALS2CR7	NM_139158	10	protein serine/threonine kinase activity	ND	ND
ANLN	NM_018685	2	actin binding	overexpressed in several tumors [4]	ND
CFH	NM_000186	5	innate immune response	overexpressed in ovarian carcinomas [5]	ND
CNOT7	NM_054026	2	transcription factor activity	ND	ND
CXCL11	NM_005409	2	chemokine activity	induction of CTL cells migration to tumors and apoptosis of tumor cells [6]	seems to contribute to metastasis of B16 melanoma cells [7]
DCN	NM_133503	8	small proteoglycan, transferase activity	tumor growth suppression [8, 9]	expressed in melanoma cell lines from different stages [10]. Inhibition of TGF-β stimulated adhesion of melanoma cells to endothelium [11]
DDX1	NM_004939	2	RNA helicase activity	co-amplified with MYCN and overexpressed in neuroblastoma and retinoblastoma cell lines [12]	ND
EIF4G2	NM_001418	2	translation initiation factor activity	downregulated in bladder carcinoma [13]	ND
FAM33A	AK056473	2	no evidence	ND	ND
FAM54A	NM_138419	2	no evidence	ND	ND
FSTL5	NM_020116	3	calcium ion binding	ND	ND

<b>GNG11</b>	NM_004126	2	signal transducer activity	downregulated in splenic marginal zone lymphoma [14]	ND
<b>HMGB2</b>	NM_002129	2	DNA binding, DNA bending activity, transcription factor activity	overexpressed, although associated to a better prognosis in epithelial ovarian cancer [15]	ND
<b>KCTD3</b>	NM_016121	2	voltage-gated potassium channel activity	ND	ND
<b>MBOAT1</b>	NM_175879	4	no evidence	ND	ND
<b>MFAP4</b>	NM_002404	3	calcium ion binding, protein binding, cell adhesion	ND	ND
<b>MYCBP</b>	NM_012333	2	transcription coactivator activity	upregulated in colon carcinoma cells overexpressing LEF-1 [16]	ND
<b>NBPF14*</b>	NM_015383	2RGP, 1MET	no evidence	ND	ND
<b>NME7</b>	NM_013330	2	kinase activity	ND	ND
<b>NUCKS1</b>	NM_022731	2	kinase activity	ND	ND
<b>NUP160</b>	NM_015231	3	nucleocytoplasmic transporter activity	ND	ND
<b>OR51E2</b>	NM_030774	2	olfactory receptor activity	overexpressed in prostate cancer [17]	ND
<b>PHF20</b>	NM_016436	2	nucleic acid binding, transcriptional regulation	overexpressed in small cell lung cancer [18]	ND
<b>POGZ</b>	NM_015100	2	nucleic acid binding, transcriptional regulation	ND	ND
<b>POLD3</b>	D26018	2	delta DNA polymerase activity	ND	ND
<b>PPT1</b>	NM_000310	2	hydrolase activity	overexpressed in colorectal carcinoma [19]	ND
<b>PSMA4</b>	NM_002789	3	threonine endopeptidase activity	ND	ND
<b>RBM25</b>	NM_021239	2	mRNA binding, mRNA splicing	ND	ND
<b>RGS2</b>	NM_002923	2	GTPase activator activity, calmodulin binding, signal transducer activity	repressed in myeloid transformation [20]	ND
<b>RUNX2</b>	NM_001015051	2	transcription factor activity	There are evidences showing runx proteins may act under different circumstances in both an oncogenic and tumor suppressor role [21]	ND
<b>SET</b>	NM_003011	2	histone binding, fosfatase inhibitor	histone acetyltransferase inhibitor, up-regulated in several tumors [22]	ND

<b>SLC35B1</b>	NM_005827	2	UDP-galactose transporter activity	ND	ND
<b>TMTC3</b>	NM_181783	2RGP, 1MET	ND	ND	ND
<b>WDR35</b>	NM_001006657	2	ND	ND	ND

See references in the reference file of the supplementary material

ND: Not described

**Table S4: Genes represented by a redundant number of sequences in the Met SSH library**

Symbol	Accession number	Redundancy	GO function	Reported in cancer	Reported in melanoma
A2M	NM_000014	12	protein carrier activity	hepatocarcinoma [23]	ND
APOD	NM_001647	2	lipid transporter activity	overexpressed in breast and prostate cancer [24]	the percentage of Apo-D positive tumors is higher in VGP than in RGP melanomas [25]
C18orf19	NM_152352	3	ND	ND	ND
CD200	NM_005944	3	integral to plasma membrane	suppression of tumor growth inhibition [26]	ND
CD59	NM_000611	9	GPI anchor binding/immune response	protects tumor cells from complement mediated lysis [27]	overexpression of CD59 in melanoma cells protects them from complement mediated lysis [28]
CHI3L2	NM_004000	2	hydrolase activity/carbohydrate metabolism	ND	ND
CPM	NM_001005502	2	GPI anchor binding, carboxypeptidase A activity, ferric iron binding , metal ion binding, metallopeptidase activity, zinc ion binding	overexpressed in tumors [29]	ND
CTSK	NM_000396	2	cathepsin activity/proteolysis	associated to tumor invasiveness [30]	ND
DCT	NM_001922	7	dopachrome isomerase activity, oxidoreductase activity/ melanin metabolism	increases proliferation of neuronal cells [31]	melanocyte/melanoma marker; increased in radioresistant melanoma cells [32]
G3BP2	NM_012297	3	RNA binding, receptor signaling complex scaffold activity	overexpressed in breast cancer [33]	ND
GPM6B	NM_001001994	3	molecular function unknown, cell differentiation	ND	ND
HLA-DRA	NM_019111	55	receptor activity, antigen presentation	observed as down-regulated[34] as also as up-regulated in different tumors [35]	prognostic significance of HLA-DR expression in melanoma tumors is not clearly defined [36, 37]
ITGB8	NM_002214	4	receptor activity, cell-matrix adhesion	overexpressed in glioblastoma [38]	ND
ITPR1	NM_002222	3	ion channel activity	ND	ND

<b>LAMA4</b>	NM_002290	2	extracellular matrix constituent, cell adhesion	angiogenesis induction [39]	ND
<b>LOC285628</b>	AL389942	3	unknown	ND	ND
<b>MLANA</b>	NM_005511	2	melanin biosynthesis (non GO)	ND	indicated as melanoma micrometastasis marker [40]
<b>MRPL42</b>	NM_172178	2	structural constituent of ribosome	ND	ND
<b>MTUS1</b>	NM_020749	3	receptor activity	considered tumor suppressor; down-regulated in pancreatic tumor [41]	ND
<b>NRP2</b>	NM_003872	2	receptor activity	expressed in gliomas and neuroblastomas [42]	expressed in melanoma [42]
<b>PDGFRA</b>	NM_006206	2	tyrosine kinase receptor	metastatic potential of oncogenic mammary epithelial cells requires an autocrine PDGF/PDGFR loop [43]	overexpressed in melanoma cells [44]
<b>PLP1</b>	NM_000533	25	structural molecule activity; involved in myelination	overexpressed in leiomyomas [45]	ND
<b>POMP</b>	NM_015932	2	protein folding, immune response	interacting partners of the human papillomavirus 16 transcription/replication factor E2 [46]	ND
<b>RDX</b>	NM_002906	2	actin binding, cytoskeletal anchoring	up-regulated in renal cell carcinoma [47]	ND
<b>SERPINE2</b>	NM_006216	4	serine-type endopeptidase inhibitor activity	overexpressed in pancreatic tumor[48]	ND
<b>SGK</b>	NM_005627	2	protein serine/threonine kinase activity	overexpressed in extraskeletal myxoid chondrosarcomas [49]	ND
<b>SLC5A4</b>	NM_014227	2	symporter activity	ND	ND
<b>SPP1</b>	NM_000582	2	cytokine activity, growth factor activity	involved in tumor progression and metastasis [50]	overexpressed in PTEN mutant melanomas [51]
<b>TDO2</b>	NM_005651	8	tryptophan 2,3-dioxygenase activity	overexpressed in esophageal adenocarcinoma [52]	ND
<b>TM4SF1</b>	NM_014220	10	molecular function unknown, integral to membrane	may be involved in cancer invasion and metastasis [53]	ND
<b>TYR</b>	NM_000372	2	monooxygenase activity; melanin biosynthesis	ND	used as molecular marker for detection circulating melanoma cells [54]

TYRP1	NM_000550	2	monooxygenase activity; melanin biosynthesis	ND	decreases the tyrosinase mediated cell death [55]
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See references in the reference file of the supplementary material

ND: Not described

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