Supplementary Information

Amigo2-upregulation in Tumour Cells Facilitates Their Attachment to Liver Endothelial Cells Resulting in Liver Metastases

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Assessment of Amigo family mRNA expression in LV12 and QRsP-11 cells

The mRNA expression levels of *Amigo1*, *Amigo2* and *Amigo3* in LV12 and QRsP-11 cells were evaluated using qRT-PCR. Bar graphs show means \pm SD (n = 4 in each group)

Supplementary Figure S2.



mRNA expression profiles of Amigo family members in HSE and LE-1 cells

Levels of *Amigo1*, *Amigo2* and *Amigo3* mRNAs in HSE and LE-1 cells were measured using qRT-PCR. Bar graphs show means \pm SD (n = 4 in each group).

Supplementary Figure S3.



Amigo2 was not required for LV12 cell adhesion to LE-1 lung endothelial cells

LV12 cells were transfected with *Amigo2* or control siRNA and their adhesion to LE-1 lung endothelial cells was then assayed. Bar graphs show means \pm SD (n = 5 in each group).

Supplementary Figure S4.



Adhesion properties of tumour cells to HSE liver endothelial cells were not changed by introduction of a luciferase expression vector

The adhesion properties of LV12-Luc or QRsP-11-Luc cells to HSE liver endothelial cells were assessed using an *in vitro* adhesion assay. Bar graphs show means \pm SD (n = 10 in each group).

Supplementary Figure S5.



Whole western blots from Figure 5a in the manuscript

The top panel is the whole gel image for western blot using anti-Amigo2 antibody in Figure 5a. The middle panel is the whole gel image for western blot using anti- β -

actin antibody in Figure 5a. The bottom panel is 0.1% Amido Black 10B (015-02192, Wako, Osaka, Japan) staining for confirmation of equivalence of the loading protein in each lane. The mouse monoclonal anti-Amigo2 antibody is described in the manufacturer's website (https://www.scbt.com/scbt/product/amigo2-antibody-g-7?productCanUrl=amigo2-antibody-g-7&_requestid=916644) and datasheet (https://datasheets.scbt.com/sc-373699.pdf).

Supplementary Table S1.

Changes in mean survival in days, liver-to-body weight ratios and Amigo2 expression following in vivo selection of a subline with high liver-metastatic properties

	No. of mice	Mean survival	Liver-to-body	Relative Amigo2
Cell line ^a	examined	in days	weight ratios (%)	mRNA expression ^b
QRsP-11	5	17.0 ± 4.0	8.4 ± 4.8	1.0 ± 0.2
LV1	5	18.4 ± 3.4	6.5 ± 3.4	3.0 ± 0.4^{d}
LV2	3	16.0 ± 0.0	20.6 ± 16.4	$3.1\pm0.2^{\text{d}}$
LV3	4	12.8 ± 2.9	6.9 ± 1.2	1.7 ± 0.4
LV4	3	21.0 ± 2.6	7.0 ± 2.2	$9.4\pm0.6^{\text{d}}$
LV5	3	20.3 ± 2.3	20.6 ± 19.0	18.0 ± 1.1^{d}
LV6	5	21.4 ± 5.2	10.9 ± 8.9	15.7 ± 1.0^{d}
LV7	4	20.3 ± 0.5	22.8 ± 20.6	30.6 ± 3.2^{d}
LV8	3	20.3 ± 3.2	32.4 ± 2.3^d	17.1 ± 0.7^{d}
LV9	5	18.4 ± 1.3	14.7 ± 13.0	17.2 ± 0.8^{d}
LV10	3	18.7 ± 1.2	30.7 ± 21.9	15.3 ± 0.8^{d}
LV11	3	18.7 ± 2.3	$28.4 \pm 11.4^{\text{c}}$	21.1 ± 2.1^{d}
LV12	7	$13.3\pm1.6^{\rm c}$	28.2 ± 9.6^{d}	19.1 ± 0.3^{d}

- a. QRsP-11 cells (1×10^6) were injected into mouse spleens. The mice were sacrificed when they appeared moribund. Tumour cells from liver metastases were cultured and used for sequential selection cycles.
- b. Amigo2 mRNA levels are indicated as fold expression relative to QRsP-11 cells.
- c. *p*<0.05 versus QRsP-11 cells.
- d. *p*<0.01 versus QRsP-11 cells.

Supplementary Table S2.

Classification of cell adhesion-related molecules upregulated or downregulated in LV12 cells compared with QRsP-11 cells

Upregulated gene		Downregulated gene	
	Fold		Fold
Name	change ^a	Name	change ^a
Integrin-related gene			
Integrin beta-like 1 (Itgbl1)	16.5	Integrin, alpha E, epithelial-	0.3
Integrin beta 7 (Itgb7)	4.3	associated (Itgae)	
Immunoglobulin-related gene	•		
Immunoglobulin superfamily,	3.6	Activated leukocyte cell	0.1
member 4A (Igsf4a)		adhesion molecule (Alcam)	
Testicular cell adhesion	2.9	Biregional cell adhesion	0.4
molecule 1 (Tcam1)		molecule-related/down-	
Poliovirus receptor (Pvr)	2.7	regulated by oncogenes	
CD4 antigen (Cd4)	2.3	(Cdon) binding protein (Boc)	
		F11 receptor (F11r)	0.4
		Coxsackievirus and	0.4
		adenovirus receptor (Cxadr)	
		Neural cell adhesion	0.5
		molecule 1 (Ncam1)	
		Neighbor of Punc E11 (Nope)	0.5
		CD47 antigen (Cd47)	0.5
Cadherin-related gene			
Protocadherin 9 (Pcdh9)	4.4	Protocadherin beta 14	0.2
		(Pcdhb14)	
		FAT tumor suppressor	0.2
		homolog 1 (Drosophila)	
		(Fat1)	
		Protocadherin beta 20	0.2
		(Pcdhb20)	
		Cadherin-like 26 (Cdh26)	0.4
		Protocadherin beta 17	0.4

		(Pcdhb17)	
		Protocadherin beta 22	0.5
		(Pcdhb22)	
Others			
Amphoterin-induced gene	15.1	Procollagen, type VI, alpha 1	0.1
and open reading frame 2		(Col6a1)	
(Amigo2)		Palladin, cytoskeletal	0.1
Wnt-1-inducible signalling	7.4	associated protein (Palld)	
pathway protein 1 (Wisp1)		Endothelial differentiation	0.1
Nexilin (Nexn)	5.0	sphingolipid G-protein-	
Contactin associated protein-	3.3	coupled receptor 1 (Edg1)	
like 4 (Cntnap4)		Neuropilin 1 (Nrp1)	0.2
Unknown	3.0	Spondin 2, extracellular	0.2
Procollagen, type VI, alpha 3	2.9	matrix protein (Spon2)	
(Col6a3)		Unknown	0.2
Pinin (Pnn)	2.8	Fibronectin 1 (Fn1)	0.2
Killer cell lectin-like receptor,	2.7	Plakophilin 2 (Pkp2)	0.2
subfamily A, member 18		Procollagen, type VI, alpha 2	0.3
(Klra18)		(Col6a2)	
Phosphatidylinositol 3-kinase,	2.4	Fibronectin leucine rich	0.3
regulatory subunit,		transmembrane protein 2	
polypeptide 1 (p85 alpha)		(Flrt2)	
(Pik3r1)		Rho family GTPase 3 (Rnd3)	0.3
Transglutaminase 2, C	2.4	B-cell leukemia/lymphoma 6	0.3
polypeptide (Tgm2)		(Bcl6)	
Thrombospondin 4 (Thbs4)	2.3	Beta-1,4-N-acetyl-	0.3
Claudin 3 (Cldn3)	2.2	galactosaminyl transferase 2	
Killer cell lectin-like receptor	2.2	(B4galnt2)	
subfamily A, member 14		Chemokine (C-X3-C motif)	0.3
(Klra14)		ligand 1 (Cx3cl1)	
G protein-coupled receptor 98	2.1	Embryonal Fyn-associated	0.4
(Gpr98)		substrate (Efs)	
Connective tissue growth	2.0	Elastin microfibril interfacer	0.4
factor (Ctgf)		2 (Emilin2)	
Kit ligand (Kitl)	2.0	von Willebrand factor A	0.4
		domain containing 1 (Vwa1)	

Procollagen, type V, alpha 2	0.5
(Col5a2)	
LIM domain containing	0.5
preferred translocation	
partner in lipoma (Lpp)	
Scavenger receptor class F,	0.5
member 2 (Scarf2)	
Nidogen 1 (Nid1)	0.5
Procollagen, type V, alpha 1	0.5
(Col5a1)	
Unknown	0.5
Thrombospondin 1 (Thbs1)	0.5

a. Cell adhesion-related genes whose expression levels had changed more than 2-fold or less than 0.5-fold in LV12 cells compared with QRsP-11 cells are indicated.

Supplementary Table S3.

Target	Strand	Sequence
Amigo1	Forward	5'-GAATGCCTTTGAGGACATGG-3'
	Reverse	5'-CTTCAGCTTGTTGGAGGACAG-3'
Amigo2	Forward	5'-GAGGCGACCATAATGTCGTT-3'
	Reverse	5'-CGGGCACCTTAGATAGGTTTT-3'
Amigo3	Forward	5'-CAGCCTCTCAGGATGGTAGC-3'
	Reverse	5'-CAGGGTGGTAAAGCCTGTGT-3'
Itgb11	Forward	5'-AGCAACCAGATGTGCAAGAA-3'
	Reverse	5'-CCATGGCCTCCACATACTTC-3'
Wisp1	Forward	5'-TTCTGCAAGTGGCCATAGGT-3'
	Reverse	5'-ACTGGCCAGGGACTCTCAC-3'
β-actin	Forward	5'-AGAGGGAAATCGTGCGTGAC-3'
	Reverse	5'-CAATAGTGATGACCTGGCCGT-3'

Primer sequences used in this study