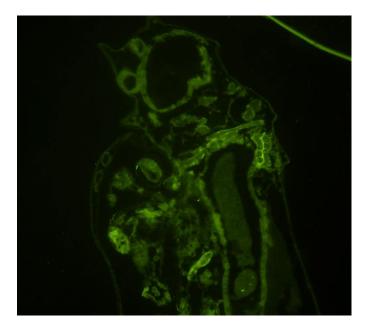
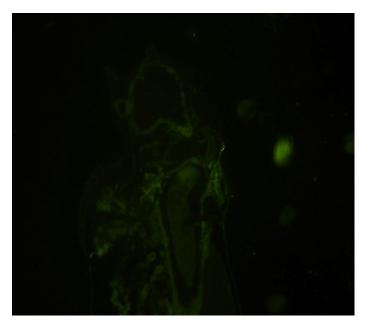
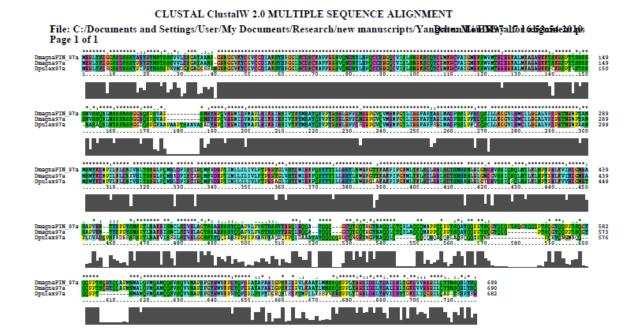
Additional File 2: Antibody recognition of HR97g is outcompeted by 100X HR97g peptide in 7-day old *Daphnia magna*



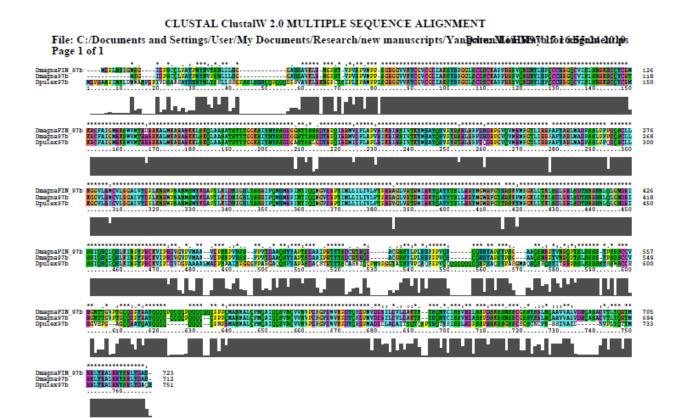
7day-D.magna-HR97g Ab



7day-D.magna-HR97g Ab + 1 to 100 dilution with HR97g peptide Additional File 3: Clustalx alignment file comparing the sequences of FinmagnaHR97a, Clem-magnaHR97a (Dmagna97a), and DappuHR97a.



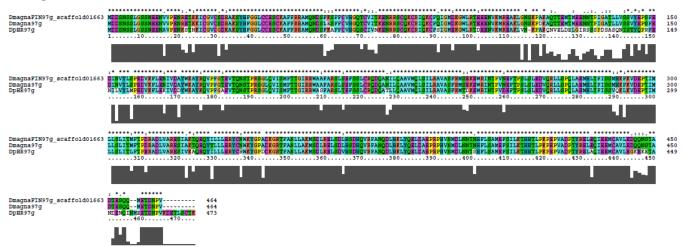
Additional File 4: Clustalx alignment file comparing the sequences of FinmagnaHR97b, Clem-magnaHR97b (Dmagna97b), and DappuHR97b.



Additional File 5: Clustalx alignment file comparing the sequences of FinmagnaHR97g, Clem-magnaHR97g, and DappuHR97g.

CLUSTAL ClustalW 2.0 MULTIPLE SEQUENCE ALIGNMENT

File: C:/Documents and Settings/User/My Documents/Research/new manuscripts/YangettenNLi/HLR97g1firb6eff3s10128lignment.ps Page 1 of 1



Additional File 6: Base-contact residues for selected NR1I, NR1J, and NR1L

members. Each member of the NR1J group has the conserved ESCKAFFR basecontact residues as does HR97g, a member of the NR1L group. HR97a and HR97b show slight variation from the NR1J conserved base-contact residue sequence.

magnaHR97g	(NR1L3)	CC <mark>ESCKAFFR</mark> RAMQND
magnaHR97a	(NR1L1)	SCD <mark>SCKAFFR</mark> RSVQND
magnaHR97b	(NR1L2)	SCD <mark>SCKAFFR</mark> RSVQND
IsHR97	(NR1L)	SCD <mark>SCKAFFR</mark> RSVQNE
magnaHR96	(NR1H)	TC <mark>ESCKAFFR</mark> RNALKG
IsHR96	(NR1H)	TC <mark>ESCKAFFR</mark> RNAIKN
DHR96	(NR1H)	TC <mark>ESCKAFFR</mark> NALAK
CeNHR48	(NR1H)	TC <mark>ESCKAFFR</mark> RNANKE
CionaVDRL	(NR1I1)	TC <mark>E</mark> G <mark>CK</mark> G <mark>FFR</mark> RSVKNS
HSCAR	(NR1I3)	TC <mark>E</mark> G <mark>CK</mark> G <mark>FFR</mark> RTVSKS
HsPXR	(NR1I2)	TC <mark>E</mark> G <mark>CK</mark> G <mark>FFR</mark> RAMKRN