

Additional File 2: Table S2. De novo assembly of the transcriptome from the seal visual cortex.

	Assembly statistics
Reads	12,473,522
Contigs	85,821
Singletons	1,919
>1000 nt	23,481
N50	1,470
Max	22,647
Average	1,061
Used for BLASTx (>500 nt)	60,602
blastx Swissprot (E<1e-5)	23,224
blastx human Refseq (E<1e-5)	22,134