

# Supplemental information #1: A new strategy to characterize the domain architecture structure of proteins of the innate immune system in tunicate species

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May 3, 2019

## Genomic sources

Sub-phylum	Specie	Source	Version
V	<i>Latimeria chalumnae</i>	Ensembl FTP <sup>1</sup>	Release 81
V	<i>Danio rerio</i>	Ensembl FTP <sup>2</sup>	Release 81
V	<i>Petromyzon marinus</i>	Ensembl FTP <sup>3</sup>	Release 81
T	<i>Ciona savignyi</i>	Ensembl FTP <sup>4</sup>	Release 81
T	<i>Ciona robusta</i>	Ensembl FTP <sup>5</sup>	Release 81
T	<i>Didemnum vexillum</i>	Universidad Nacional de Colombia <sup>8</sup>	v.2
T	<i>Botryllus schlosseri</i>	ANISEED <sup>9</sup>	v.1
T	<i>Botrylloides leachii</i>	ANISEED <sup>10</sup>	v.1
T	<i>Molgula occidentalis</i>	ANISEED <sup>11</sup>	v.1
T	<i>Molgula oculata</i>	ANISEED <sup>12</sup>	v.1
T	<i>Oikopleura dioica</i>	OikoBase <sup>13</sup>	Version 3
C	<i>Branchiostoma floridae</i>	JGI genome portal <sup>14</sup>	v.1 and v.2
H	<i>Saccoglossus kowalevshii</i>	NCBI FTP <sup>15</sup>	Skow_1.1
E	<i>Patiria miniata</i>	Echinobase <sup>16</sup>	v2.0
E	<i>Strongylocentrotus purpuratus</i>	Echinobase <sup>17</sup>	v4.2

Table 1: Genomic data source. Described labels for Subphylums: **V**: vertebrates, **T**: tunicates, **C**: cephalochordates, **H**: hemichordates and **E**: echinoderms.

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<sup>1</sup>[ftp://ftp.ensembl.org/pub/release-81/fasta/latimeria\\_chalumnae/pep/Latimeria\\_chalumnae.LatChal1.pep.all.fa.gz](ftp://ftp.ensembl.org/pub/release-81/fasta/latimeria_chalumnae/pep/Latimeria_chalumnae.LatChal1.pep.all.fa.gz)

<sup>2</sup>[ftp://ftp.ensembl.org/pub/release-81/fasta/danio\\_rerio/pep/Danio\\_rerio.GRCz10.pep.all.fa.gz](ftp://ftp.ensembl.org/pub/release-81/fasta/danio_rerio/pep/Danio_rerio.GRCz10.pep.all.fa.gz)

<sup>3</sup>[ftp://ftp.ensembl.org/pub/release-81/fasta/petromyzon\\_marinus/dna/Petromyzon\\_marinus.Pmarinus\\_7.0.dna.toplevel.fa.gz](ftp://ftp.ensembl.org/pub/release-81/fasta/petromyzon_marinus/dna/Petromyzon_marinus.Pmarinus_7.0.dna.toplevel.fa.gz)

<sup>4</sup>[ftp://ftp.ensembl.org/pub/release-81/fasta/ciona\\_savignyi/pep/Ciona\\_savignyi.CSAV2.0.pep.all.fa.gz](ftp://ftp.ensembl.org/pub/release-81/fasta/ciona_savignyi/pep/Ciona_savignyi.CSAV2.0.pep.all.fa.gz)

<sup>5</sup>[ftp://ftp.ensembl.org/pub/release-81/fasta/ciona\\_intestinalis/pep/Ciona\\_intestinalis.KH.pep.all.fa.gz](ftp://ftp.ensembl.org/pub/release-81/fasta/ciona_intestinalis/pep/Ciona_intestinalis.KH.pep.all.fa.gz)

<sup>6</sup><http://tunicatadvexillum.bioinf.uni-leipzig.de/>

<sup>7</sup>[https://www.aniseed.cnrs.fr/aniseed/download/?file=data%2Fbs%2Fbotryllus\\_protein\\_fasta.zip](https://www.aniseed.cnrs.fr/aniseed/download/?file=data%2Fbs%2Fbotryllus_protein_fasta.zip)

<sup>8</sup>[https://www.aniseed.cnrs.fr/aniseed/download/?file=data%2Fboleac%2FBoleac\\_proteins\\_v4\\_fasta.zip](https://www.aniseed.cnrs.fr/aniseed/download/?file=data%2Fboleac%2FBoleac_proteins_v4_fasta.zip)

<sup>9</sup>[https://www.aniseed.cnrs.fr/aniseed/download/?file=data%2Fmoocci%2Fmolgula\\_occidentalis\\_protein\\_fasta.zip](https://www.aniseed.cnrs.fr/aniseed/download/?file=data%2Fmoocci%2Fmolgula_occidentalis_protein_fasta.zip)

<sup>10</sup>[https://www.aniseed.cnrs.fr/aniseed/download/?file=data%2Fmoocul%2Fmolgula\\_oculata\\_protein\\_fasta.zip](https://www.aniseed.cnrs.fr/aniseed/download/?file=data%2Fmoocul%2Fmolgula_oculata_protein_fasta.zip)

<sup>11</sup><http://oikoarrays.biology.uiowa.edu/Oiko/Downloads.html>

<sup>12</sup><https://genome.jgi.doe.gov/Brafl1/Brafl1.home.html>

<sup>13</sup>[ftp://ftp.ncbi.nlm.nih.gov/genomes/Saccoglossus\\_kowalevskii/protein/protein.fa.gz](ftp://ftp.ncbi.nlm.nih.gov/genomes/Saccoglossus_kowalevskii/protein/protein.fa.gz)

<sup>14</sup>[http://www.echinobase.org/Echinobase/PmDownload/pmin\\_proteins\\_v2.0.fa](http://www.echinobase.org/Echinobase/PmDownload/pmin_proteins_v2.0.fa)

<sup>15</sup>[ftp://ftp.ncbi.nlm.nih.gov/genomes/all/GCF/000/002/235/GCF\\_000002235.4\\_Spur\\_4.2/GCF\\_000002235.4\\_Spur\\_4.2\\_protein.faa.gz](ftp://ftp.ncbi.nlm.nih.gov/genomes/all/GCF/000/002/235/GCF_000002235.4_Spur_4.2/GCF_000002235.4_Spur_4.2_protein.faa.gz)