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Identification and tissue distribution by *in-silico* analysis and RT-qPCR expression profiling of chicken c-type lectin-like domain containing proteins



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We investigated

- Genes in the chicken genome encoding c-type lectin-like domain containing proteins (CTLDcps)
 - Gene validation by tissue expression profiling of identified CTLDcps in respiratory tissue and spleen

Why

- To find CTLDcps in the chicken genome and potential mammalian orthologues
 - Verification of mRNA expression in respiratory tissues and spleen

We found

- 15 novel chicken CTLDcps of which CLEC17AL-A and -B belong to group II C-type lectins
 - An avian specific expansion of group V C-type lectins
 - Differential tissue expression of identified CTLDcps

How

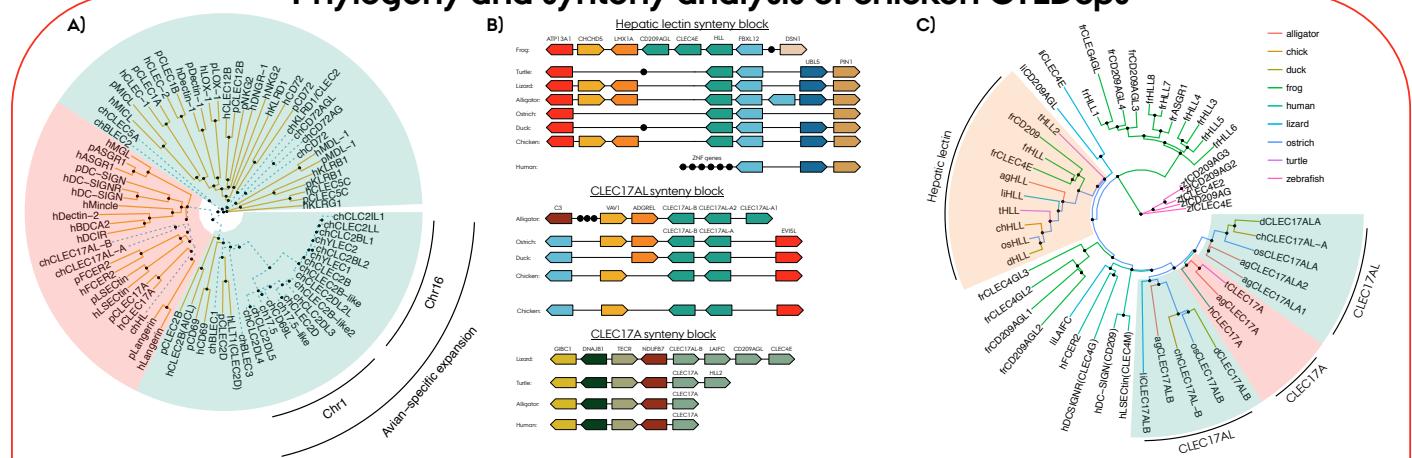
In silico analysis (GRCg6a):

- BLASTp
 - Synteny analysis
 - Phylogenetic analysis

RT-qPCR:

- B15 day-old chicks (n=3)
• Tissues: NALT Trachea
Lung Spleen

Phylogeny and synteny analysis of chicken CTLDcps



A) Phylogeny of chicken (blue dotted lines) and mammalian (yellow lines):

- *chHL*, *chCLEC17AL-A*, and -*B* belong to group II CTLDcps (**red**)
- Avian specific expansion of group V (**green**) chicken CTLDcps located on Chr16 and Chr1

- B)** Synteny for:

 - *chCLEC17AL-A* and *-B* between birds and alligator
 - *CLEC17A* in reptiles and human
 - *chHL* between reptiles and frog

C) Phylogeny of group II CTLDCPs:

- Orthologues of *chHL* present in reptiles and amphibians (**yellow**) but not in human
- Gene lineages to *chCLEC17AL-A* and -*B* on one side (**green**), and human *CLEC17A* (**red**) on the other side, evolved separately from a common ancestral gene

RT-qPCR expression profiling in respiratory tissue and spleen

Gene name	ACB symbol	Protein refseq	CDS	Genomic location
<i>CICD25</i>	CICD25L	XP_0151285725.1	1	NC_000985.2 (1888..3504)
<i>CICB28-like</i>	CICB28L	XP_0010749487.1	1	NC_000985.2 (19192..19563, complement)
<i>CICD24</i>	CICD24L	XP_0250085640.1	1	NC_000985.6 (45693..58403)
<i>D6H9</i>	D6H9L	NP_001156303.1	1	NC_000985.6 (23720..24165)
<i>CICD23</i>	CICD23L	NP_001246797.1	1	NC_000985.6 (30135..303113)
<i>GLC2C</i>	GLC2CL	XP_011747821.1	1	NC_000985.6 (217130..221977)
<i>GLC2B</i>	GLC2BL	XP_011747822.1	1	NC_000985.6 (217130..221977, Scott et al., 2012)
<i>GLCB28-like</i>	GLCB28L	XP_0250117001.1	1	NC_000985.1 (58495..197427, complement)
<i>CICB2L</i>	CICB2L	XP_015156641.1	16	NC_000985.10 (199752..200939)
<i>CICB2L</i>	CICB2L	XP_0250117579.1	16	NC_000985.10 (202530..202849)
<i>CICL21</i>	CICL21L	XP_025011683.1	16	NC_000985.10 (198831..197460, complement)
<i>GLCB2</i>	GLCB2L	XP_0250107469.1	16	NC_000985.10 (200914..209342)
<i>GLCB2-like</i>	GLCB2L	XP_0250107470.1	16	NC_000985.10 (200914..209342, complement)
<i>B6C3</i>	B6C3L	NP_001157050.1	16	NC_000985.10 (197994..197474)
<i>B6C3</i>	B6C3L	NP_001030626.1	16	NC_000985.10 (202446..244761, complement)
<i>ZOT24ZL</i>	ZOT24HL	XP_001231020.2	2	NC_000985.12 (9940..1542, 965503)
<i>CLECSA</i>	CLECSA	XP_000432020.1	1	NC_000985.12 (9940..1542, 965503)
<i>Hepsic lectin</i>	HEPSL	NP_909181.5	30	NC_000985.12 (102577..106772, 107916)
<i>CETC17AL1</i>	CETC17AL1	XP_003443419.1	30	NC_000985.29 (1952..199516)
<i>CETC17AL8</i>	CETC17A8L	XP_0237795.30	30	NC_000985.29 (9957..104620)

- | Gene name | NCBI symbol | Protein refseq | Chr | Genomic location |
|-------------|--------------|----------------|-----|---|
| CLEC2L5 | CLEC2L5 | XP_015183575.1 | 1 | NC_0040085 (21188..35004) |
| CLEC2-like | CLEC2L4 | XP_050085992.1 | 1 | NC_0050885 (19419..19583), complement |
| CLEC2L4 | CLEC2L4 | XP_050085401.1 | 1 | NC_0050885 (4693..58430) |
| CD49L | CD49L | XP_015164303.1 | 1 | NC_0040085 (23720..24165) |
| CLEC2L3 | CLEC2L3 | NP_0126497.1 | 1 | NC_0040085 (30435..30513) |
| CLEC2D | LOC101748271 | NP_015128239.1 | 1 | NC_0040085 (21710..21977) |
| CLEC2B | CLEC2B | NP_015128240.1 | 1 | NC_0040085 (21710..21977), Scott et al., 2016, complement |
| CLEC2B-like | LOC101748270 | NP_050117901.1 | 1A | NC_2001015 (19849..19972), complement |
| CLEC2B-like | CLEC2B | NP_015165643.1 | 1A | NC_2001015 (19849..20093), complement |
| CLEC2L1 | CLEC2L1 | XP_015175793.1 | 1A | NC_0040085 (203520..20849) |
| CLEC2L1 | CLEC2L1 | NP_025011683.1 | 1A | NC_0061035 (19883..19970), complement |
| CLEC2B | LOC101889209 | XP_05011773.1 | 16 | NC_0061035 (202913..20932) |
| CLEC2D-like | LOC10506497 | XP_05011773.1 | 16 | NC_0061035 (23810..24393), complement |
| GC2A | GC2L2 | XP_015167541.1 | 16 | NC_0061035 (210790..21924) |
| BLEC3 | BLEC3 | NP_013006261.1 | 16 | NC_0061035 (24840..24479), complement |
| DZT4AGE | DZT4AGE | NP_015123062.2 | 2 | NC_0040085 (21710..21977) |
| CLEC2A | CLEC2A | NP_000390031.1 | 2 | NC_0040085 (21710..21977), complement |
| HLL | HLL | NP_99015.1 | 30 | NC_028739.2 (106377..10479) |
| CLEC17A-L | FCER2 | XP_03043419.1 | 30 | NC_028739.2 (159521..159951) |
| CLEC17AL | CLEC17A | XP_403779.5 | 30 | NC_028739.2 (19975..20462) |

