

RARA

History

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%%{init: { 'logLevel': 'debug', 'theme': 'dark' } }%%
timeline
  title Publication timing
    2017-10-10 : Reddy : DLBCL
```

Relevance tier by entity

Entity	Tier	Description
<div>DLBCL</div>	2	relevance in DLBCL not firmly established

Mutation incidence in large patient cohorts (GAMBL reanalysis)

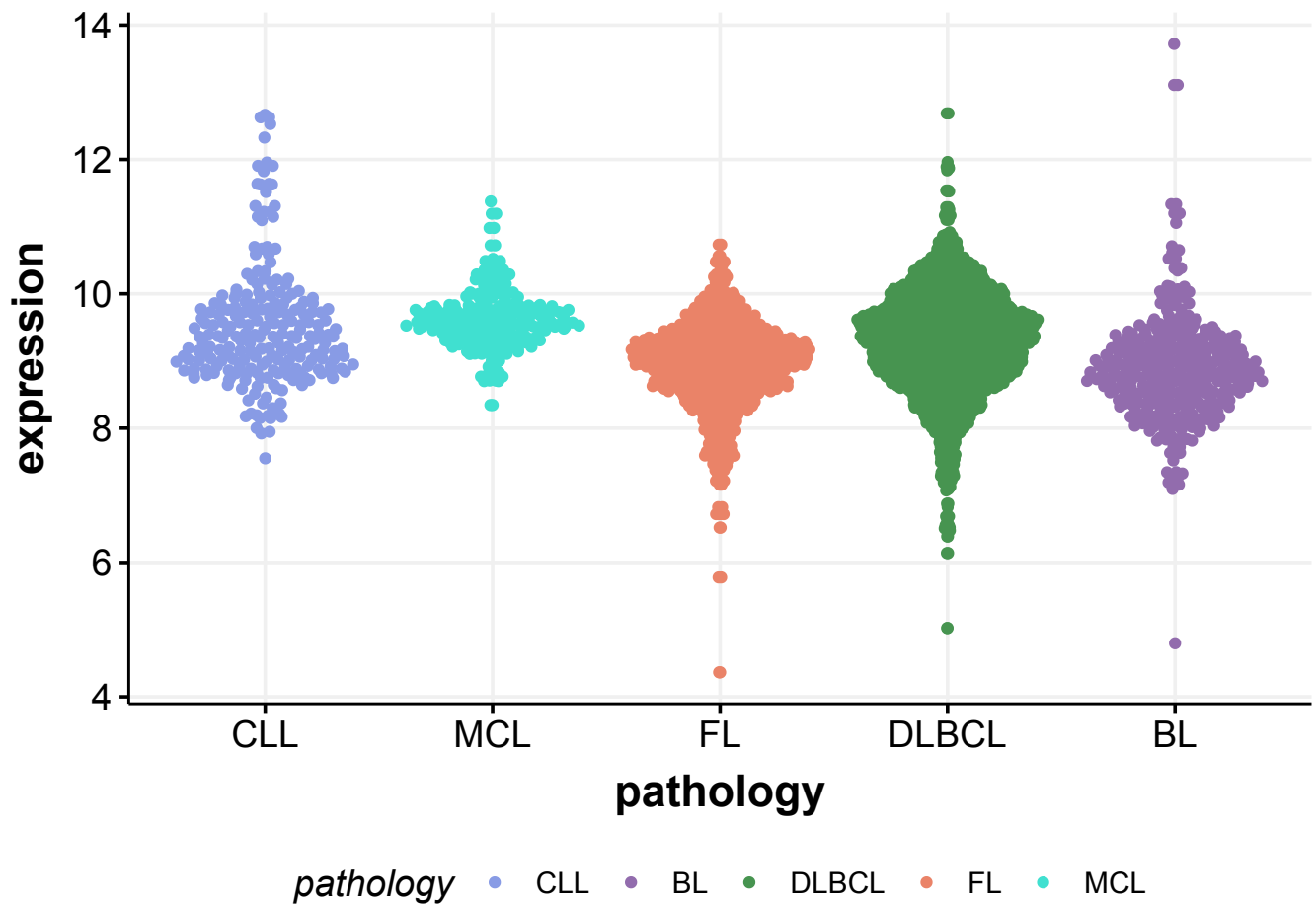
Entity	source	frequency (%)
DLBCL	GAMBL genomes	1.34
DLBCL	Schmitz cohort	4.04
DLBCL	Reddy cohort	0.70
DLBCL	Chapuy cohort	0.43

Mutation pattern and selective pressure estimates

Entity	aSHM	Significant selection	dN/dS (missense)	dN/dS (nonsense)
BL	No	No	0.000	0
DLBCL	No	No	6.208	0
FL	No	No	0.000	0

[!NOTE] First described in DLBCL in 2017 by [Reddy A](#)

View coding variants in ProteinPaint [hg19](#) or [hg38](#)



References

1. Reddy A, Zhang J, Davis NS, Moffitt AB, Love CL, Waldrop A, Leppa S, Pasanen A, Meriranta L, Karjalainen-Lindsberg ML, Nørgaard P, Pedersen M, Gang AO, Høgdaal E, Heavican TB, Lone W, Iqbal J, Qin Q, Li G, Kim SY, Healy J, Richards KL, Fedoriw Y, Bernal-Mizrachi L, Koff JL, Staton AD, Flowers CR, Paltiel O, Goldschmidt N, Calaminici M, Clear A, Gribben J, Nguyen E, Czader MB, Ondrejka SL, Collie A, Hsi ED, Tse E, Au-Yeung RKH, Kwong YL, Srivastava G, Choi WWL, Evens AM, Pilichowska M, Sengar M, Reddy N, Li S, Chadburn A, Gordon LI, Jaffe ES, Levy S, Rempel R, Tzeng T, Happ LE, Dave T, Rajagopalan D, Datta J, Dunson DB, Dave SS. Genetic and Functional Drivers of Diffuse Large B Cell Lymphoma. *Cell*. 2017 Oct;171(2):481-494.e15.