


# CCND1

## History

```
%%{init: { 'logLevel': 'debug', 'theme': 'dark' } }%%
timeline
  title Publication timing
    2013-11-05 : Bea : MCL
```

## Relevance tier by entity

Entity	Tier	Description
 MCL	1	high-confidence MCL gene[@beaLandscapeSomaticMutations2013]

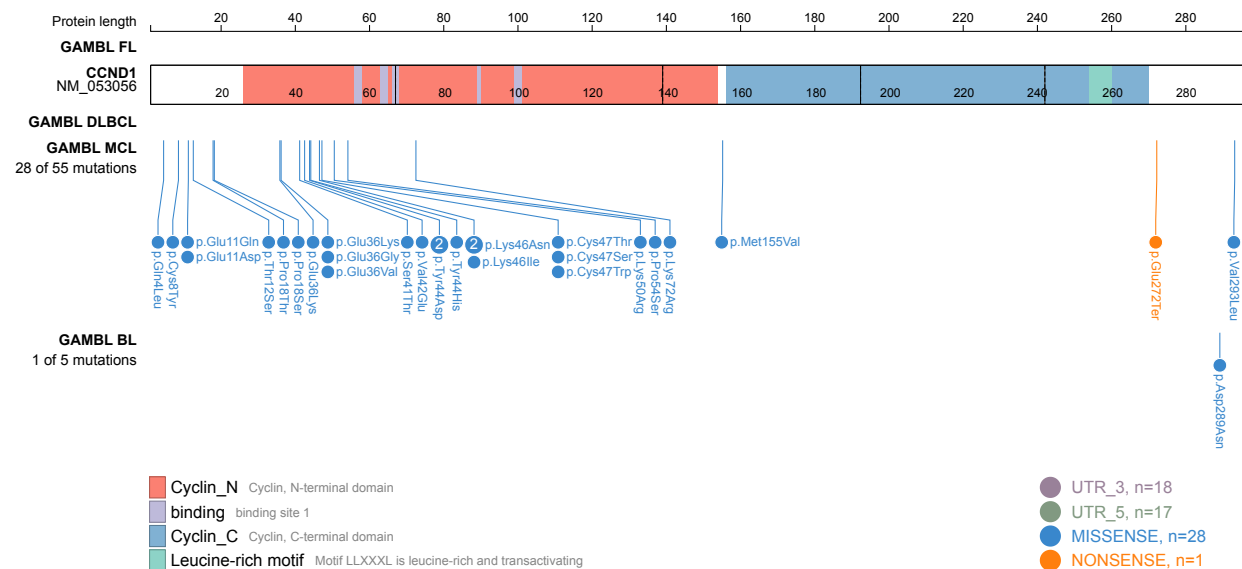
## Mutation incidence in large patient cohorts (GAMBL reanalysis)

Entity	source	frequency (%)
MCL	GAMBL genomes	14.69

## Mutation pattern and selective pressure estimates

Entity	aSHM	Significant selection	dN/dS (missense)	dN/dS (nonsense)
BL	No	No	1.909	0
DLBCL	No	No	0.000	0
FL	No	No	0.000	0

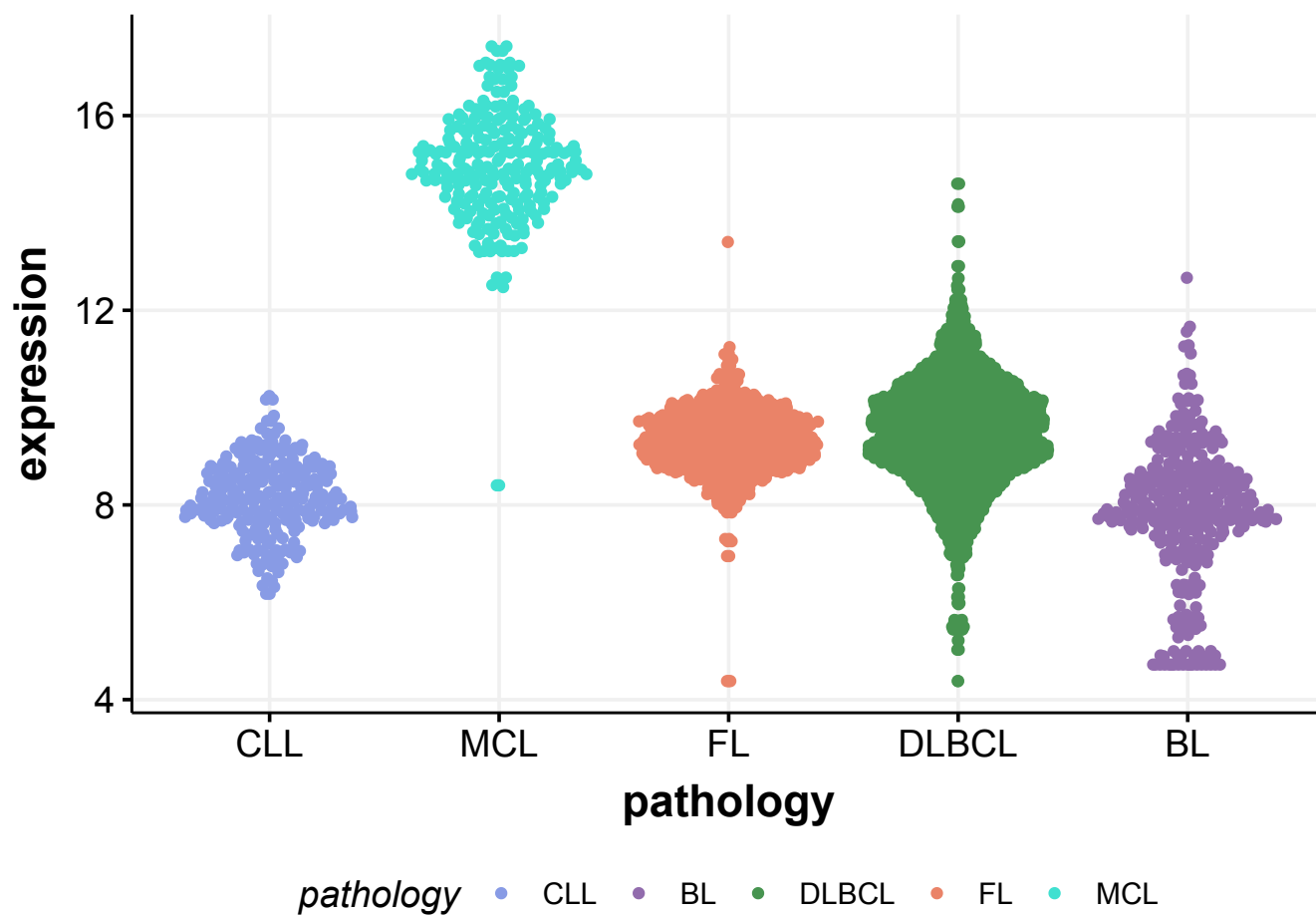
View coding variants in ProteinPaint [hg19](#) or [hg38](#)



View all variants in GenomePaint [hg19](#) or [hg38](#)



CCND1 Expression



References