

B-NHL colours

load main table with all hex codes

```
full_codes = read_tsv("colour_codes.tsv") %>% dplyr::filter(is.na(is_alias)) %>% select(-is_alias)
```

```
## Rows: 223 Columns: 5
## -- Column specification -----
## Delimiter: "\t"
## chr (5): category, group, name, colour, is_alias
##
## i Use `spec()` to retrieve the full column specification for this data.
## i Specify the column types or set `show_col_types = FALSE` to quiet this message.
```

Full palette

[illegible]

Hex codes for B-cell lymphomas

[illegible]

Cell of Origin / Gene Expression Subgroups

[illegible]

LymphGen classes with Composite classes

Name	Code	category	group
		subgroup	LymphGen
		subgroup	LymphGen
		subgroup	LymphGen
		subgroup	LymphGen
		subgroup	LymphGen
		subgroup	LymphGen
		subgroup	LymphGen
		subgroup	LymphGen
		subgroup	LymphGen
		subgroup	LymphGen
		subgroup	LymphGen
		subgroup	LymphGen
		subgroup	LymphGen
		subgroup	LymphGen
		subgroup	LymphGen

HMRN classes

Name	Code	category	group
		subgroup	hmrn
		subgroup	hmrn
		subgroup	hmrn
		subgroup	hmrn
		subgroup	hmrn
		subgroup	hmrn
		subgroup	hmrn
		subgroup	hmrn

Clinical variables

[illegible]

BL and DLBCL Genetic subgroups from the Morin lab

Name	Code	category	group
		subgroup	BL-genetic
		subgroup	BL-genetic
		subgroup	BL-genetic
		subgroup	BL-genetic
		subgroup	BL-genetic
		subgroup	BL-genetic
		subgroup	DLBCL-genetic
		subgroup	DLBCL-genetic
		subgroup	DLBCL-genetic
		subgroup	DLBCL-genetic
		subgroup	DLBCL-genetic
		subgroup	DLBCL-genetic
		subgroup	DLBCL-genetic

ASH family journals

Name	Code	category	group
		publication	blood
		publication	blood
		publication	blood
		publication	blood
		publication	blood
		publication	blood
		publication	blood
		publication	blood
		publication	blood
		publication	blood
		publication	blood
		publication	blood
		publication	blood
		publication	blood
		publication	blood