Sorting RNAi info by accession ID

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You can find RefSeq accession IDs on RNAi product pages, but start your search with a target gene/protein

Although we capture accession IDs (RefSeq) of RNAi targets/genes in our database, they are not searchable parameters at this time. Generally, our **search strategy** is to select a product type, name your target gene or protein and further refine using our filters from there.

Honing in on splice variants and specific isoforms, however, is a feature we are still developing. Though, you can try searching for the variant directly in the search bar. **Let us know** if the variant you are searching for is missing!

If searching by accession ID is something that would be valuable to your workflow, we would love to **talk to you** to learn more about it.