

LINCS Field Name	Related to	Description	Comments	Importance	Centers Provide
<b>RN Name</b>	canonical	The primary name of the siRNA or shRNA as chosen by LINCS	Should be descriptive and correspond to existing siRNA or shRNA names as much as possible; batch independent name	1	YES
<b>RN LINCS ID</b>	canonical	Unique LINCS internal identifier	LINCS internal ID; this is a batch independent ID; canonical siRNA or shRNA ID	1	-
<b>RN Probe ID</b>	canonical	ID of the siRNA or shRNA as listed in NCBI Probe database	-	2	YES
<b>RN Probe Title</b>	canonical	Name of the siRNA or shRNA as listed in NCBI Probe database	-	2	YES
<b>RN Probe Type</b>	canonical	A controlled vocabulary specifying whether the probe is an siRNA , esiRNA, or shRNA	-	1	YES
<b>RN shRNA Construct</b>	canonical	A description of the RNA construct includes the name of vector, the gene that is targeted by the siRNA/shRNA, gene ID, regulatory region, selectable marker.	for shRNA only	2	YES
<b>RN shRNA Vector Reference</b>	canonical	Reference to publication or contact information (if applicable)	for shRNA only	2	-
<b>RN Target Gene Symbol</b>	canonical	The NCBI Entrez Gene Symbol for the gene targeted by the siRNA or shRNA	It might be desirable to request the sequence of the target mRNA, especially if a particular splice variant is targeted. Also, it should be noted whether the target sequence lies in the coding region or UTR of the mRNA.	1	YES
<b>RN Target Gene ID</b>	canonical	The NCBI Entrez Gene ID for the gene targeted by the siRNA or shRNA	-	1	-
<b>RN RNAi Sense Sequence</b>	canonical	The nucleotide sequence of the sense (passenger) strand of the siRNA or the processed shRNA.	Note that several vendors do not make their siRNA sequences public. Thus, it might be impossible to require exact sequence information for all siRNAs. These vendors will submit context sequences to NCBI Probe db though, so one would always have Probe IDs.	2	-
<b>RN Validation Information</b>	canonical	Information about experimental verification of siRNA/shRNA activity. A reference (PubMed or other suitable reference) should be provided.	Information about the cell line/cell type and organism used for validation, as well as the % reduction in protein expression and mRNA observed in the validation experiments; whether the target monitored in validation studies was the endogenous mRNA or a transfected mRNA. It would be useful information even if only a code is given that states whether an RNAi reagent was validated or not by LINCS effort or by others.	2	-
<b>RN Provider Name</b>	batch	Vendor or lab that supplied the reagent	-	1	YES
<b>RN Provider Catalog ID</b>	batch	ID or catalogue number assigned to the reagent by the vendor or provider	-	1	YES
<b>RN Provider Batch ID</b>	batch	Batch or lot number assigned to the reagent by the vendor or provider	-	1	YES
<b>RN Silencing Reagents</b>	experiment	Number of combined silencing RNA reagents per well	From MIARE	1	-
<b>RN Delivery Type</b>	experiment	Describe type of delivery, e.g. reverse transfection, infection, electroporation, intravenous injection, shooting, feeding	From MIARE	1	-
<b>RN Delivery Reagent</b>	experiment	Delivery reagent description, including, type, name, catalog number, manufacturer, and final concentration	From MIARE	2	-
<b>RN Concentration Of Silencing Reagent</b>	experiment	The final concentration of the silencing reagents	From MIARE	2	-
<b>RN Cell Number</b>	experiment	Number of cells per well in the delivery plate	From MIARE	1	-
<b>RN Assay Conditions</b>	experiment	Time to assay point from delivery of silencing RNA reagent; time of exposure of silencing RNA reagent; media changes	From MIARE	2	-
<b>RN Number Of Replicates</b>	experiment	The number of replicates used in the experiment	From MIARE	1	-
<b>RN Center Name</b>	batch	LINCS center using the RNAi	-	1	YES