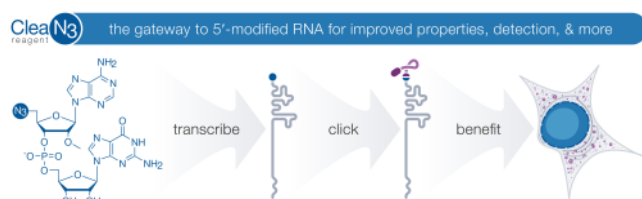


Enhancement of RNA properties via in vitro transcription priming

A new molecular tool utilising in vitro transcription priming to modify RNA, enhancing its stability and translation, and allowing its detection.

Applications: Cap-independently translated mRNA, Therapeutic vaccines, Gene therapies, Cancer immunotherapies, Regulatory control of protein expression, mRNA screening, mRNA detection, mRNA delivery



Features

Use of an azido-functionalised dinucleotide in vitro transcription primer, Clean3, that relies on base-pairing with T7 RNA polymerase promoters leads to high priming efficiencies and in vitro transcription yields

The 5' azide present in Clean3-primed transcripts, provides access to post-transcriptional modification via click chemistry, e.g., strain-promoted azide-alkyne cycloaddition (SPAAC)

Benefits

Superior in vitro transcription yields, priming efficiencies, and primed transcript homogeneities over existing primers for cap-independently translated mRNAs, and non-coding RNAs

A rapid and efficient process over existing synthesis processes – one working day to synthesis 5' modified transcripts from nucleoside triphosphates; Click chemistry permits improvements to properties of cap-independently translated mRNAs

Features	Benefits
Enhanced linear cap-independently translated (CIT) mRNA properties, such as translation efficiency and stability without eliciting immunogenicity	Enables production of therapeutic mRNA designs to target diseases, such as cancers; Spatiotemporal characterisation of CIT mRNA uptake and cellular fate
	Convenient benchmarking of internal ribosome entry sites (IRESs) and cap-independent translational enhancers (CITEs) independent of cell line-specific factors, which is currently challenging

Awaiting patent and Available for

- Co-development
- Consulting
- Licensing

To find out more, contact enquiries@innovation.ox.ac.uk and quote the project number.

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