ASH1L methyltransferase: a therapeutic target in prostate cancer?

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Background

The histone 3 lysine 36 (H3K36) when methylated is crucial for transcriptional regulation, splicing, DNA repair and genome stability [1]. The tri-methylation status of H3K36 has been shown to guide RNA methylation co-transcriptionally [2]. Both methyltransferases and demethylases that target H3K36me have been shown to play a role in cancer. ASH1L, trithorax histone methyltransferase, that di-methylates H3K36, has shown to function through binding of MRG15 at the chromatin [3] and is involved in global nuclear excision repair [4]. Recent study has shown, that compared to the other H3K36 di-methylators NSD1 and NSD2, ASH1L methylation is restricted to active regulatory elements of developmental genes [5]. ASH1L is not well studied in prostate cancer, with a recent study showing its role in castrate resistant AR-negative prostate cancer (CRPC) [6]. With the development of ASH1L inhibitor [7], this study aims to identify whether ASH1L is a potential therapeutic target in AR-positive CRPC.

Methods

Immunohistochemistry staining performed in a UK cohort of 100 tumour and 45 non-malignant samples and H-scored. AR-positive CRPC cell lines were utilised to measure mRNA (QPCR) and protein levels of ASH1L (western blotting). AS-99, ASH1L inhibitor was used to look at cell proliferation (measuring DNA content), invasion (transwell assay) and transcriptome (RNA-Seq). Combination therapies of AS-99 and enzalutamide were tested.

Results

ASH1L nuclear levels did not vary significantly in primary adenocarcinomas compared to nonmalignant samples. ASH1L levels varied in AR-positive CRPC cell lines and was no longer androgenregulated. ASH1L methyltransferase inhibition resulted in, differential gene expression, splicing, loss of proliferation and invasion. Inhibition of ASH1L lead to sensitisation of 22RV1 cells to enzalutamide.

Conclusions: ASH1L could be a potential target for inhibition in AR-positive CRPC to resensitise to current AR-targeted therapies. Ongoing investigation is underway on pinpointing ASH1L target genes and ASH1L regulation in CRPC.

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