

## Study identifies new drug target in deadly form of leukemia

June 3 2013

A research team led by the Duke-NUS Graduate Medical School (Duke-NUS) in Singapore has identified ways to inhibit the function of a key protein linked to stem cell-like behavior in terminal-stage chronic myeloid leukemia (CML), making it possible to develop drugs that may extend the survival of these patients.

The study, published in the prestigious international journal *Proceedings* of the National Academy of Sciences, is the result of a long-standing collaboration between Duke-NUS, the Experimental Therapeutics Centre at the Agency for Science, Technology and Research (A\*STAR), and the Singapore General Hospital that is focused on developing effective therapies in CML.

CML is a <u>blood cancer</u> that has seen tremendous improvement in treatment outcomes following the introduction of <u>tyrosine kinase</u> <u>inhibitor</u> (TKI) drugs that specifically target the BCR-ABL <u>fusion gene</u>, a <u>genetic abnormality</u> that is characteristic of CML. However, when CML progresses to its terminal stage, known as the <u>blast crisis</u> phase, TKI drugs become ineffective and patients with blast crisis CML rapidly succumb to the disease.

"TKI therapy is highly effective in chronic phase CML, and enables most patients to survive many years. In contrast, patients with blast crisis CML usually succumb to their disease within one year, with most patients dying because they develop drug resistance to TKI therapy," said principal investigator Ong Sin Tiong, associate professor and head of the



Laboratory of <u>Hematologic Malignancies</u> in the Cancer and <u>Stem Cell Biology</u> Program at Duke-NUS.

A subset of cells associated with blast crisis CML exhibit characteristics of self-renewing stem cells, suggesting that targeting this particularly malignant and drug-resistant population would be effective in treating blast crisis CML. The team therefore searched for novel targets that will specifically eliminate these cancer stem cells.

Through their efforts, the team identified a protein enzyme, known as the MNK kinase, that was abnormally activated in clinical samples taken from patients with blast crisis CML. Experiments conducted in the lab further unraveled how MNK kinase activation plays a critical role in the progression of CML to the blast crisis phase, and confers stem cell-like behavior on blast crisis cells.

The team tested a panel of drugs that inhibit MNK kinase activity and found that these MNK inhibitors were effective in preventing blast crisis cells from behaving like cancer stem cells in both in vitro laboratory tests and animal studies.

"Our studies identify the MNK kinases as an important therapeutic target in blast crisis CML, and suggest that drug inhibition of MNK kinase will be useful in overcoming TKI resistance, and improving the survival of patients with blast crisis CML," said Ong, who is also a visiting consultant at the National Cancer Center Singapore and Singapore General Hospital.

Importantly, the MNK inhibitor drugs do not appear to be toxic to normal blood stem cells, indicating that drugs targeting MNK kinases may not cause harmful side effects. Ong said he hopes the findings from this study will open new research directions in the treatment of blast crisis CML.



"We are currently collaborating with the Experimental Therapeutics Centre and Singapore General Hospital to develop new drugs to simultaneously target the MNK and the BCR-ABL kinases. The development of dual MNK and BCR-ABL kinase inhibitors to treat patients with blast crisis CML may enhance the survival of patients with this deadly disease," Ong said. He added he ho estimates it will take a few years before these drugs can enter into clinical trials for blast crisis CML.

**More information:** Targeting of the MNK–eIF4E axis in blast crisis chronic myeloid leukemia inhibits leukemia stem cell function, <a href="https://www.pnas.org/cgi/doi/10.1073/pnas.1301838110">www.pnas.org/cgi/doi/10.1073/pnas.1301838110</a>

## Provided by Duke University Medical Center

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