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(fig. S5A)
reveal a certain
degree of
mobility of the
[6P4]CCL5 core,
the receptor ...

Structural basis

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activation of
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receptor 5 by a
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agonist
Co-Investigator
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as a model, we
measured the
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terminal phospho-
acceptor sites
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3 (S3), and Ser
7 (S7) to
various stimuli,
including GPCR
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modeling suggest
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methylation of
peptides
enhances their
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nitrogen atoms
within peptide
bonds

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will be
probabilistic,
and the study
will be carried
out through
computer
simulation. The
required
background is
knowledge of

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random processes
(commonly used
discrete ...

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may have future
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proteome of
stromal tissue
in an
individualized
but coordinated
manner
These bind to
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binding proteins
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(bacterial
enzymes that
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the analysis,

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design of

sewerage systems

have been

available since

the 1970s.

Complex models

are now readily

available. The

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models includes
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