Meeting abstract

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The sorting protein GASP-I regulates the constitutive signaling capacity of the virally encoded chemokine receptor US28 Pia Tschische¹, Elisabeth Moser¹, Dawn Thompson², Wolfgang Platzer¹, Henry Vischer³, Martine J Smit³, Helmut Schaider⁴, Lene Martini², Jennifer Whistler² and Maria Waldhoer^{*1}

Address: ¹Institute of Experimental and Clinical Pharmacology, Medical University of Graz, 8010 Graz, Austria , ²Ernest Gallo Clinic and Research Center, University of California San Francisco, CA 94608, USA, ³Leiden/Amsterdam Center for Drug Research (LACDR), Division of Medicinal Chemistry, Faculty of Sciences, VU University Amsterdam, 1081 HV Amsterdam, The Netherlands and ⁴Department of Dermatology, Medical University of Graz, 8036 Graz, Austria

Email: Maria Waldhoer* - maria.waldhoer@medunigraz.at

* Corresponding author

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Background

Human cytomegalovirus (HCMV) is a widespread pathogen that has been shown to be present in various malignancies and it is also thought to be linked to vascular diseases. HCMV encodes the seven transmembrane (7 TM)/G protein-coupled receptor (GPCR) US28, which constitutively activates the $G\alpha_q$ /phospholipase C (PLC) pathway and downstream transcription factors such as the nuclear factor- κ B (NF- κ B) or the cyclic AMP responsive element binding protein (CREB). In this study we set out to elucidate the role of the GPCR-associated sorting protein-1 (GASP-1) in the regulation of the constitutive signaling capacity of US28.

Methods

To elucidate the role of GASP-1 in the regulation of the constitutive signaling capacity of US28 we disrupted the US28/GASP-1 interaction by either overexpression of dominant negative cGASP-1 or shRNA knock-down of endogenous GASP-1. To monitor the US28-mediated signaling we conducted inositol phosphate (IP) accumulation assays as well as luciferase reporter gene assays to check the activation of the transcription factors NF- κ B and CREB.

Results

We find that GASP-1 is indeed able to modulate the signaling activity of US28. Disruption of the GASP-1/US28 interaction by either i) overexpression of dominant negative cGASP-1 or by ii) shRNA knock-down of endogenous GASP-1 alters the US28-mediated $G\alpha_q$ /PLC/IP accumulation as well as the activation of the transcription factors NF- κ B and CREB.

Conclusion

By identifying the sorting protein GASP-1 as a key regulator of the constitutive signaling activity of US28, we may be one step closer to gaining a better understanding of this viral receptor and its significance in the pathogenesis implicated by HCMV.