### **AUG 2**

# *Supplementary Information for*

# **Synthesis and Characterization of Heparosan-Granulocyte-Colony Stimulating Factor Conjugates: a natural sugar-based drug delivery system to treat neutropenia**

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### **Supplementary Table**



### **Supplementary Table 1: Summary of hematological changes in rats injected with G-CSF conjugates**.

The fold-increase is listed above for various blood cell types at various times post-injection in comparison to the buffer vehicle alone. (*n* = ~5/group; values represents change from concurrent control on the given study period time).

### **Supplementary Figure 1. Structure of Acceptor and HEP derivatives**

First, the heparosan polymer was produced by synchronized stoichiometrically controlled polymerization using a heparosan trisaccharide amine-containing glycoside acceptor (**A**) and UDP-GlcA and UDP-GlcNAc with the HEP synthase. This HEP-NH<sub>2</sub> polymer ( $n = \sim 145$  or  $\sim 260$  for the 55- or 99-kDa polymers, respectively) was purified. In the second step,  $HEP-NH<sub>2</sub>$  was converted to the aldehyde form via reaction with a heterobifunctional reagent to create HEP-CHO (**B**).

The HEP-CHO and G-CSF were coupled by reductive amination at pH 5 to more selectively react with the amino terminus of the protein (rather than lysine side-chains) yielding the HEP-G-CSF conjugate (**C**). To define the coupling site, the conjugate was first treated with heparin lyase III (cleaves the sugar chain, leaving an unsaturated glucuronic acid derivative at the non-reducing end) and the protein with a 3-sugar unit 'stub' was PAGE purified. Second, the relevant gel slice was excised and treated with the protease trypsin to generate an oligosaccharide-peptide species (**D**) that was identified by liquid chromatography/mass spectrometry (*see* **Supp. Fig. 5**).

# **Supplementary Fig. 1**



 $R_1$  = G-CSF  $R_2 =$  amino-terminal tryptic peptide



#### **Supplementary Figure 2. Electrophoretic analyses of 99-kDa HEP-G-CSF and strong anion exchange (SAX) chromatography purification of 55-kDa HEP-GCSF.**

**Panel** *a* - Samples were separated in 4-20% TGX gradient gel and stained with Coomassie Blue. **std**, Precision Plus® protein size standard. **Lane 1**, free G-CSF. **Lane 2**, conjugation reaction mixture before purification. **Lane 3**, uncoupled protein in the flow-through of SAX. **Lane 4**, blank. **Lane 5**, purified HEP-G-CSF conjugate (marked with a *star*).

**Panel b** - SAX profile of the 55-kDa-HEP-G-CSF purification depicted in Fig. 2 of main text. The peak marked with the *arrow* was pooled and used for animal efficacy testing (*blue*, UV absorbance at 215 nm; *orange*, NaCl gradient).

**std G P-G**



# **Supplementary Fig 3. SDS-PAGE analysis of purified PEG-G-CSF**.

The starting protein and the in-house prepared, cation exchange chromatography-purified PEGconjugate were analyzed on 4-20% TGX gel with Coomassie Blue staining. **Lanes: G**, G-CSF; **P-G**, PEG-G-CSF (*star*); **std**, Precision Plus® protein standard.



**log [protein] (ng/ml)**

# **Supplementary Fig 4. NFS-60 cell proliferation assay of 55-kDa and 99-kDa HEP-G-CSF and PEG-G-CSF.**

All 3 conjugates stimulated proliferation in culture. The approximate effective concentration for 50% effect (EC<sub>50</sub>) values (pg/ml) were: PEG-G-CSF, 4.1; 55-kDa HEP, 20; 99-kDa HEP, 25.

#### **Intensity**



### **Supplementary Figure 5. LC-MS-MS analysis of trisaccharide-peptide 'stub' of HEP-G-CSF conjugate.**

The 55-kDa HEP modified G-CSF (*orange*) or unmodified protein (*blue; grey* normalized to HEP-G-CSF intensity for other peptide species) were analyzed in parallel by peptide mapping. A unique +4-charged species (composed of MTPLGPASSLPQSFLLK:  $C_{82}H_{136}N_{19}O_{23}S_1$  and Hep3linker:  $C_{30}H_{40}N_2O_{18}$  forming the complex:  $C_{112}H_{174}N_{21}O_{41}S_1$ ; standard formula weight: 2502.761; *see* Supp. Fig. 1, structure **D**) was observed confirming coupling of the heparosan chain to the amino-terminus of the protein drug.

### **Supp. Fig 6. Molecular simulation data for hydrodynamic radius predictions of HEP-G-CSF conjugates**.

The data below for the 20-kDa, 55-kDa, or 99-kDa HEP-G-CSF was used to generate the values in **Table I**; the *green trace* reflects the average  $r_h$  over time (cumulative moving average). Note that the longer sugar chains displayed a wider range of possible configurations and consequently the same number of computations resulted in a less accurate average  $r_h$  prediction.



# 20-kDa HEP-G-CSF



# 55-kDa HEP-G-CSF

**Supp. Fig 6. Molecular simulation data for hydrodynamic radius predictions of HEP-G-CSF conjugates**. (cont.)





### **Supplementary Figure 7. Hematology study with single injection of HEP-G-CSF (55- or 99 kDa), PEG-G-CSF (in-house prepared), or Neulasta® in healthy rats.**

The absolute white blood cell counts were stimulated by all conjugates in comparison to the vehicle control (*black*).



### **Supplementary Figure 8**. **Immunization and bleeding schedule of the long-term immunological study in three Sprague Dawley rats.**

The rats were boosted with 55-kDa HEP-G-CSF antigen nine times approximately every three weeks. *Arrows* indicate the time of antigen boost (*black*) and bleed collection (*gray*). Each rat received 55-kDa HEP-G-CSF about every three weeks at 0.4-0.7 mg/kg. Serum were collected before the first injection (week zero = 'prebleed') and at week 9, 15 and 27 (rat #3) or at week 9, 15 and 30 (rat #1, and #2) (post-injection bleed 1, 2, and 3). To detect potential anti-heparosan antibodies, enzyme-linked immunosorbent assay (ELISA) plates were coated with HEP-BSA (a conjugate with bovine serum albumin). As a negative control, some wells were treated with BSA only. The presence of any potential anti-HEP-G-CSF antibodies was analyzed by ELISA with goat anti-rat IgG or IgM as the secondary antibodies (as horseradish peroxidase conjugates; *see* Fig. 5 for results).