

RDP1258, a New Rationally Designed Immunosuppressive Peptide, Prolongs Allograft Survival in Rats: Analysis of Its Mechanism of Action

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Abstract

Peptides derived from the HLA class I heavy chain (a.a. 75–84) have been shown to modulate immune responses in vitro and in vivo in a non-allele-restricted fashion. In vivo studies in rodents have demonstrated prolonged allograft survival following peptide therapy. The immunomodulatory effect of these peptides has been correlated with peptide-mediated modulation of heme oxygenase 1 activity (HO-1). Recently, we used a rational approach for designing novel peptides with enhanced immunosuppressant activity. These peptides were also more potent inhibitors of HO-1 activity in vitro. Here we evaluated one of these peptides, RDP1258, for its ability to prolong heterotopic heart graft survival in rats. The peptide mediated effect on HO-1 was analyzed in vitro and in vivo. Peptide RDP1258 was shown to inhibit rat HO-1 in vitro in a dose-dependent fashion. However, RDP1258, like other HO-inhibitors,

when administered to rats, secondarily resulted in an up-regulation of splenic HO-1 activity. Up-regulation of HO-1 was associated with prolonged heart allograft survival (6.6 ± 0.6 vs. $2/14 > 100$ days and $12/14$ 16.2 ± 1.7 days; $p < 0.001$). The analysis of graft infiltrating cells on day 5 after transplantation showed a significant decrease in the number of graft infiltrating cells in RDP1258-treated recipients compared to untreated ones (14.8 vs. 32.7% ; $p < 0.01$). In addition, grafts from peptide-treated animals showed significantly decreased expression of TNF- α mRNA and increased levels of iNOS mRNA. Our results are consistent with the recent observation that up-regulation of HO-1 results in the inhibition of several immune effector functions. Modulation of HO-1 activity may enable the development of novel immunomodulatory strategies in humans.

Introduction

The main molecular targets of the alloimmune response are the major histocompatibility complex (MHC) molecules. Over the past decade,

several investigators have evaluated the use of soluble MHC molecules to modulate immune responses and allograft rejection. Synthetic peptides corresponding to various regions of MHC molecules have been shown to affect T cell functions in vitro. Krensky and others (1–3) showed that synthetic peptides derived from the class I heavy chain inhibited lysis by cytotoxic T lymphocytes (CTLs). In particular, two peptides corresponding to amino acid 75–84 of the $\alpha 1$ helix

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of HLA B-2702 (2702.75-84) and B-0701 (07.75-84) have been shown to inhibit the differentiation of cytotoxic T cells *in vitro* in a non-allele-restricted fashion (i.e., the effect of the peptides is the same, regardless of donor haplotype or target specificity) (4,5). In addition to their effect on T cells, these peptides also decreased natural killer (NK) cell-mediated cytotoxicity, suggesting that their mechanism of action is independent of the indirect presentation that has been shown to be responsible for the biological activity of other allele-specific HLA-derived peptides. Peptides 2702.75-84 and 07.75-84 prolonged allograft survival in rodents and induced tolerance when administered in conjunction with subtherapeutic doses of cyclosporin A (CsA) (6–9). We have recently shown that these peptides' immunomodulatory activity is correlated with peptide-mediated modulation of the activity of heme oxygenase 1 (HO-1), also called HSP32 (10). Novel peptides with enhanced immunomodulatory activity have been developed by computer-assisted rational design based on *in vivo* data (11). These peptides have been shown to be efficient inhibitors of HO-1 *in vitro*. In the present study, one of these peptides, RDP1258, was assessed for its ability to modulate immune responses in rats and to affect heterotopic heart graft survival. Peptide RDP1258 was found to prolong allograft survival significantly, and this prolonged survival was associated with enhanced expression of splenic HO-1. Moreover, allografts from treated recipients showed a significant decrease in the number of graft-infiltrating cells, particularly in the CD68/monocyte population; decreased expression of tumor necrosis factor α (TNF- α) mRNA; and an increased level of inducible nitric oxide synthase (iNOS) mRNA. These results support the hypothesis that RDP1258-mediated immunomodulation involves HO-1 activity and monocyte/macrophage functions.

Materials and Methods

Reagents

Peptides were synthesized using F-moc chemistry, purified by HPLC, and shown to be >90% homogeneous by analytical reverse phase HPLC (SynPep, Dublin, CA). The following peptides were used: 2702.75-84, RENLRALRY; 07.75-84, RESLRNLRGY; and RDP1258, RNleNleNleRNleNleNleGY-CONH₂, where Nle is norleucine. Metalloporphyrins, purchased from Porphyrin Products (Logan, UT), were dissolved in a small

quantity of 0.2 M NaOH, subsequently adjusted to pH 7.4 with 1 M HCl and diluted in 0.85% NaCl. The stock concentration of metalloporphyrins was 1 mg/ml.

Antibodies

The following mouse monoclonal antibodies (MAbs) were used to phenotype rat leukocytes: OX19 (anti-CD5, T cells), ED1 (recognizing most tissue macrophages, and monocytes, granulocytes, and dendritic cells), MRC OX33 (anti-CD45 isoform present on B cells only), MRC OX1 (anti-CD45), MRC OX8 (anti-CD8), and W3/25 (anti-CD4). All these MAbs were purchased from ECACC. The mouse anti-rat MAb 3.2.3, reacting with NK cells, was generously donated by W. H. Chambers (Pittsburgh, PA). For HO-1 staining, a polyclonal rabbit anti-rat HO-1 antibody was used (OSA100 Sress-gen), followed by a peroxidase-conjugated chicken anti-rabbit IgG (Vector Laboratories).

HO-1 Enzyme Assay

To determine HO activity in tissues, rat spleen samples were homogenized on ice in a Tris-HCl lysis buffer (pH 7.4) containing 0.5% Triton X-100 and protease inhibitors. Samples were frozen in small aliquots until use. Spleen homogenate (100 μ l) was mixed with 0.8 mM NADPH, 0.8 mM glucose-6-phosphate, 1.0 U G-6-P dehydrogenase, 1 mM MgCl₂, and 10 μ l purified rat liver biliverdin reductase at 4°C. The reaction was initiated by the addition of hemin (final concentration, 0.25 mM). The reaction mixture was incubated at 37°C in the dark for 15 min. At the end of the incubation period, any insoluble material was removed by centrifugation and supernatants were analyzed for bilirubin concentration by measuring the difference in absorbance at 460 and 530 nm. An extinction coefficient of 40 mM⁻¹ cm⁻¹ at A 460–530 was used to calculate the amount of bilirubin formed. Controls included spleen samples in the absence of the NADPH generating system, and all the ingredients of the reaction mixture in the absence of spleen homogenates. Biliverdin reductase was purified from rat liver using the method described by Kutty and Maines (12).

Heart Allograft Transplantation

Eight- to 12-week-old male Lewis.1W (RT1.^u) and Lewis.1A (RT1.^a) rats served as donors and

recipients of heart allografts, respectively. These congenic strains were obtained from the Centre d'Élevage Janvier (Le Genest-Saint-Isle, France). Heterotopic heart grafts were performed using Ono and Lindsey's technique (13). The grafts were evaluated daily for function by palpation through the abdominal wall. Lewis.1A recipients rejected Lewis.1W heart allografts in 6 ± 0.7 days.

Immunohistology

Tissue samples for immunohistology were embedded in Tissue Tek (OCT compound), snap frozen in liquid nitrogen immediately after harvesting, and stored at -70°C until used. Five micrometer cryostat sections were cut at -20°C , air dried, fixed in acetone for 10 min at room temperature, and stored at -20°C . Tissue sections were labeled using a three-step indirect immunoperoxidase technique (6). Negative controls were performed, omitting the first antibody.

Quantitative Analysis of Cellular Infiltrate

The area of each immunoperoxidase-labeled tissue section infiltrated by leukocytes of a particular phenotype was determined by morphometric analysis using the point-counting technique (6). With the aid of a 121 intersection-bearing square grid in the eyepiece, sections were examined at a magnification of $400\times$ by two observers. For each of 15 adjacent high-power fields, the number of positively stained cells superimposed by an intersection was counted and the percentage area of each section occupied by cells of a particular phenotype was calculated as follows: [(number of positive cells under grid intersections)/(total number of grid intersections)] \times 100.

Sera

Blood samples were obtained by heart puncture under ether anesthesia and clotted at 37°C . Sera were heat inactivated at 56°C for 30 min, and stored at -20°C .

Preparation of Cell Suspensions

Infiltrating mononuclear cells were mechanically harvested from heart allografts without the use of enzymes, as previously described (6). Briefly, heart grafts were finely minced and gently pressed through a stainless steel mesh, filtered

twice, and layered on a Ficoll-Hypaque gradient. Splenocytes were flushed by multiple intra-splenic injections of phosphate-buffered saline (PBS) and mononuclear cells were obtained after a Ficoll-Hypaque gradient density.

Detection of Anti-Donor Alloantibodies

Concanavalin-A (Con A) blasts were prepared from LEW.1W. Con-A splenocytes, 1×10^5 in $30 \mu\text{l}$, were incubated with $30 \mu\text{l}$ diluted sera (1:2, 1:4, 1:8 and 1:16) for 45 min at 4°C , and washed twice in PBS containing 0.5% bovine serum albumin (BSA) (Sigma) and 0.02% sodium azide (Az) (PBS-BSA-Az). To stain for IgG alloantibodies, washed cells were made to react with $30 \mu\text{l}$ of PBS-BSA-Az containing fluorescein isothiocyanate (FITC) conjugated F (ab')² goat antibody specific for the Fc portion of rat IgG, (Jackson Immunoresearch Labs, West Grove, PA) for 30 min at 4°C . After two additional washes, cells were fixed in 1% neutral buffered formalin and analyzed on a FACS calibur (Becton-Dickinson, Mountain View, CA).

Cytotoxic Assays

Graft infiltrating cells from treated and untreated recipients were tested for alloantigen-specific and nonspecific cytotoxicity using a standard 6 hr ⁵¹Cr-release assay as described elsewhere (6). Allo-specific cytotoxicity was assessed using Con-A blasts from LEW.1W (RT1^u targets) and LEW.1A (RT1^a targets). Spontaneous cytotoxicity was assessed using the NK-susceptible YAC cell line. Specific ⁵¹Cr-release was calculated according to the formula $100 \times (\text{experimental release} - \text{spontaneous release}) / (\text{maximum release} - \text{spontaneous release})$. Data are shown as the mean of triplicate determinations of 25:1 ratio, spontaneous release being $<20\%$ of maximum release in all experiments.

Competitive Quantitative PCR

The sequences of the oligonucleotides are listed in Table 1. Oligonucleotides designated as X.Cons encompass the X.5' primer, four bases of the X cDNA sequence, followed by a deletion of four bases and ending with the corresponding sequence of the X cDNA sequence (14). Total RNA from heart allografts was purified in a cesium chloride gradient (15). Ten micrograms was reverse transcribed, using the Boehringer cDNA synthesis kit, and diluted in a final volume of 100

Table 1. Oligonucleotide sequences

Primers	Sequences (5'-3')
iNOS-5'	ACTCAAGTTCAGCTTGGCGG
iNOS-3'	GGAGTGCAG TGGCTCCAG
iNOS RO	CATGCGGCCT CCTTGAGC
TNF- α -5'	CTACTGCTTCAGCTCCACAG
TNF- α -3'	GACCCGTAGGGCGATTACAG
TNF- α -RO	GCTTCCCAACGCTGGGTCC

μ l. Standard constructions were performed essentially as previously described (14). Quantification of transcript X was performed as previously described (14). Briefly, a constant amount of cDNA corresponding to the reverse transcription of 50 ng of total RNA was mixed with 10^6 , 10^5 , 10^4 , 10^3 , 10^2 , or 0 copies of the X standard, and then amplified to saturation. This was followed by a dye-labeled oligonucleotide elongation (runoff reaction), and the runoff reaction products were electrophoresed for 7 hr using an Applied Biosystem 373A DNA Sequencer. The Immunoscope[®] software was specially devised to measure both the length of each DNA peak detected (with a precision of <0.2 nucleotide) and its area (with a precision of <5%) (14).

Statistical Analysis

To compare the number of mRNA copies and the percentage of antibody labeling in each of the two experimental groups, the Student's *t*-test was performed, and $p < 0.05$ was considered significant. Allograft survival times were compared using the Kaplan-Meier test.

Results

Peptide RDP1258-Mediated Modulation of Rat HO-1 Activity

Recently, peptides 2702.75-84, 07.75-84 and derivatives thereof have been shown to inhibit HO-1 activity in vitro (10). The rational design of peptide RDP1258 was based solely on in vivo data (prolongation of heart allograft survival) (11). To determine whether peptide RDP1258 interacted with rat HO-1, enzyme activity in spleen homogenates was measured in vitro in the presence of increasing concentrations of pep-

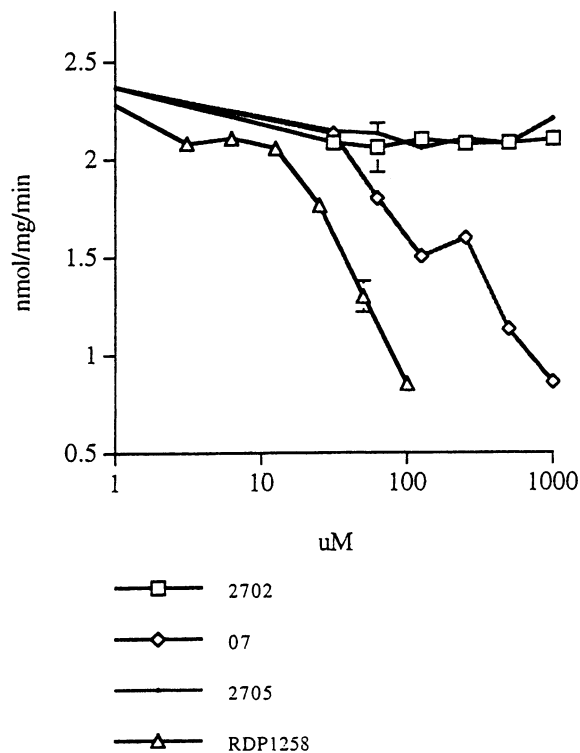


Fig. 1. Effect of peptide on in vitro HO activity. The activity of heme oxygenase in a spleen cell extract was measured in the presence of increasing amounts of peptide 2702.75-84 (squares), 07.75-84 (diamonds), 2705.75-84 (line), and RDP1258 (triangles). Enzyme activity is expressed as nmol bilirubin formed/mg of protein/min.

ptide (Fig. 1). Half-maximal inhibition of HO-1 activity by peptide 07.75-84 (IC_{50}) was observed at 200 μ M. Peptide RDP1258 appeared to be more effective, the observed IC_{50} being 20 μ M. Peptide 2702.75-84 had no effect on rat HO-1 activity, which is consistent with the observation that it had no effect on rat allograft survival (6,16). In contrast, peptide 07.75-84, which was shown to be effective in rats but not in mice, inhibited rat HO-1 activity (6,16). Moreover, the results show that in vitro, RDP1258 peptide inhibited HO-1 activity in a dose-dependent fashion.

It is well established that inhibition of HO-1 in vivo results in a rapid up-regulation of HO-1 production (17-19). The effect of peptide RDP1258 on rat HO-1 was evaluated by measuring splenic HO-1 activity at various time points following administration of the peptide (Fig. 2). Injection of 0.5, 1.0, and 2.0 mg/kg resulted in increases of 22%, 66%, and 45%, respectively, in splenic HO-1 activity 24 hr after administra-

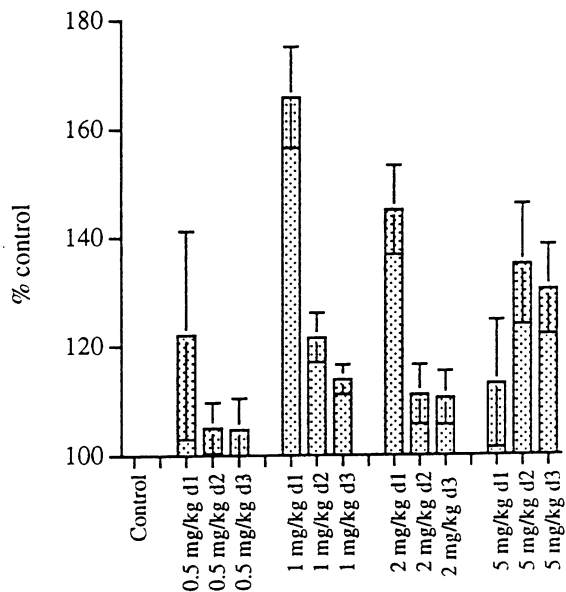


Fig. 2. Effect of peptide on splenic HO activity after in vivo treatment. Splenic HO activity was measured 24, 48, and 72 hr following injection of the indicated amounts of peptide RDP1258. HO activity in peptide-treated animals ($n = 3$) was compared to mannitol-treated animals ($n = 3$) and is expressed as % of control. HO activity in mannitol-treated control animals was 3.081 ± 0.051 nmol/mg/min.

tion; this activity then declined to almost normal levels within the next 24 hr. Following the injection of 5 mg/kg, the increase observed after 24 hr was 13%. After 48 and 72 hr, increases of 35% and 30%, respectively were observed. This delayed but sustained increase in HO-1 activity may be due to the prolonged continuing pres-

ence of peptide RDP1258 following injection of this high dose (5 mg/kg).

The expression of splenic HO-1 was also evaluated in allograft recipients by immunohistochemistry. Splens from untreated and RDP1258-treated allograft recipients (four animals per group) were removed 5 days after transplantation and labeled with anti-HO-1 polyclonal antibody (Fig. 3). Splens from untreated recipients (Fig. 3A) showed almost no staining for HO-1 protein. In splens from RDP1258-treated recipients (Fig. 3B), the HO-1 staining was intense in particular in the "marginal" zone of the spleen, which contains mostly monocytes and macrophages. We also analyzed the expression of HO-1 protein in heart allografts by immunoperoxidase staining. There is an increased expression of HO-1 in allografts compared to naive heart, but we did not find any significant difference between allografts from both groups (data not shown).

Prolonged Allograft Survival following Peptide RDP1258 Therapy

The effect of peptide RDP1258 therapy on allograft survival was investigated in LEW.1A recipients of LEW.1W heart allografts. Recipients were treated before transplantation with different doses of the peptide. Heart allografts in untreated recipients were rejected in 6.6 ± 0.6 days (Fig. 4 and Table 2). Peptide RDP1258 therapy at 1 mg/kg on days -7 and -1 significantly prolonged allograft survival (2/14 > 100 days and 12/14 16.2 ± 1.7 days vs. 6.6 ± 0.6 in the untreated group; $p \leq 0.001$). No further effect was observed when the dose was increased to 5

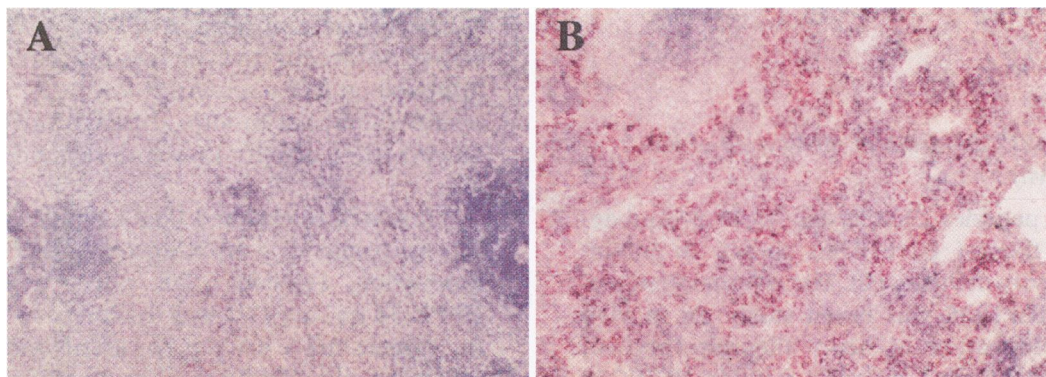


Fig. 3. HO-1 protein expression in spleen of allograft recipients. Immunostaining with an anti-HO-1-specific polyclonal antibody in spleen from untreated (A, 20 \times amplification) and RDP1258-treated (B, 20 \times amplification) allograft recipients 5 days after transplantation.

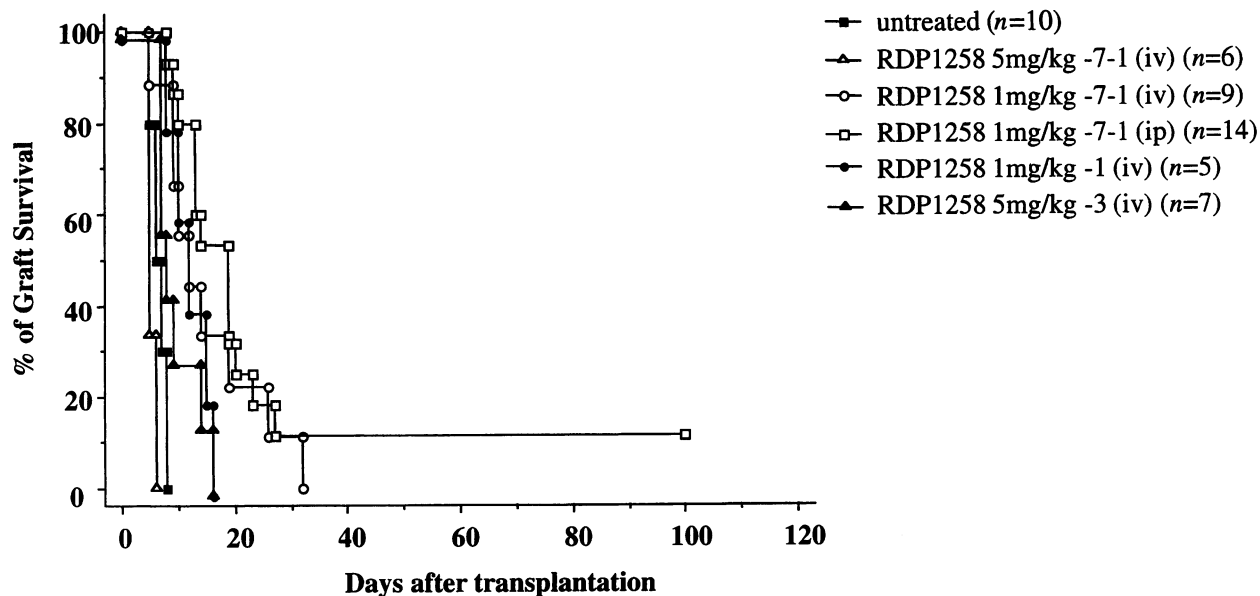


Fig. 4. Allograft survival after RDP1258 therapy. Kaplan-Meier depiction of heart allograft survival in rats receiving RDP1258 peptide therapy.

mg/kg on days -7 and -1. Interestingly, when the peptide was injected on day -3 at 5 mg/kg, there was a small but significant ($p < 0.009$) prolongation of allograft survival compared to the effect of the same dose given on days -7 and -1 (Table 3). These results are in agreement with the effect (dose and timing) of peptide therapy in HO-1 activity (Fig. 2). A comparison of intravenous (IV) and intraperitoneal (IP) administration suggested that IP administration is as effective as IV administration (in the IP group 2/14 > 100 days and in 12/14 the mean allograft

survival was 16.2 ± 1.7 days vs. 15.1 ± 2.9 in the IV group).

ZnPP Abrogates the Effect on Allograft Survival of Peptide RDP1258 Therapy

Zinc-protoporphyrin (ZnPP) is a well-known inhibitor of HO-1 activity (20–23). To test if the effect of RDP1258 peptide was mediated in part by an increase in HO-1 activity, the administration of ZnPP was investigated to see whether it inhibits the effect of RDP1258 treatment. Recipients were treated with RDP1258 on days -7 and -1 IP at 1 mg/kg and the day of the graft they were treated with ZnPP at 10 mg/Kg subcutaneously (SC). As shown in Table 3, recipients treated with ZnPP rejected their graft in 7.2 ± 0.8 days, suggesting that the effect of RDP1258 therapy was cancelled by ZnPP treatment ($p < 0.02$).

Prolonged Allograft Survival following Cobalt-protoporphyrin Therapy

Cobalt-protoporphyrin (CoPP) is a known inhibitor of HO-1 (17–19). The administration of CoPP was therefore investigated to see whether it might mimic some of the peptide’s effect. It was shown to result in up-regulation of HO-1 similar to that observed with peptide RDP1258. In addition, CoPP has been shown to prolong heart allograft survival in mice (24). We therefore studied its effect on allograft survival in rats.

Table 2. IgG alloantibody production in sera of allograft recipients 5 and 7 days after transplantation

	IgG	
	Day 5	Day 7
Untreated (n = 4)	142 ± 32	185 ± 0.5
RDP1258-treated (n = 4)	156 ± 28	225 ± 2.5

Sera samples obtained from untreated and RDP1258-treated allograft recipients at 5 and 7 days after transplantation were tested for IgG anti-donor specific alloantibody production against donor cells (see Materials and Methods). Results are expressed as the mean of channel fluorescence of four samples per group.

Table 3. RDP1258 peptide effects on allograft survival

Treatment	Graft survival	Mean \pm SEM
No treatment	5, 5, 6, 6, 6, 7, 7, 8, 8, 8	6.6 \pm 0.6
RDP1258 5 mg/kg i.v. day -7, -1	5, 5, 5, 5, 6, 6	5.3 \pm 0.2
RDP1258 5 mg/kg i.v. day -3	7, 7, 7, 8, 9, 14, 16	9.7 \pm 1.4 ^a
RDP1258 1 mg/kg i.v. day -7, -1	5, 9, 9, 10, 12, 14, 19, 26, 32	15.1 \pm 2.9 ^b
RDP1258 1 mg/kg i.v. day -1	8, 10, 12, 15, 16	12.2 \pm 1
RDP1258 1 mg/kg i.p. day -7, -1	8, 9, 10, 13, 13, 14, 19, 19, 19, 20, 23, 27, >100, >100	16.2 \pm 1.7 ^c
RDP1258 1 mg/kg i.p. day -7, -1 and ZnPP 20 mg/kg s.c. day 0	6, 6, 6, 6, 8, 11	7.2 \pm 0.8 ^d
ZnPP 20 mg/kg s.c. day 0	7, 8, 13, 14	10.5 \pm 1.8

^a $p < 0.009$ compared to RDP1258 5 mg/kg IV day -7, -1 group.

^b $p < 0.0035$ compared to untreated group.

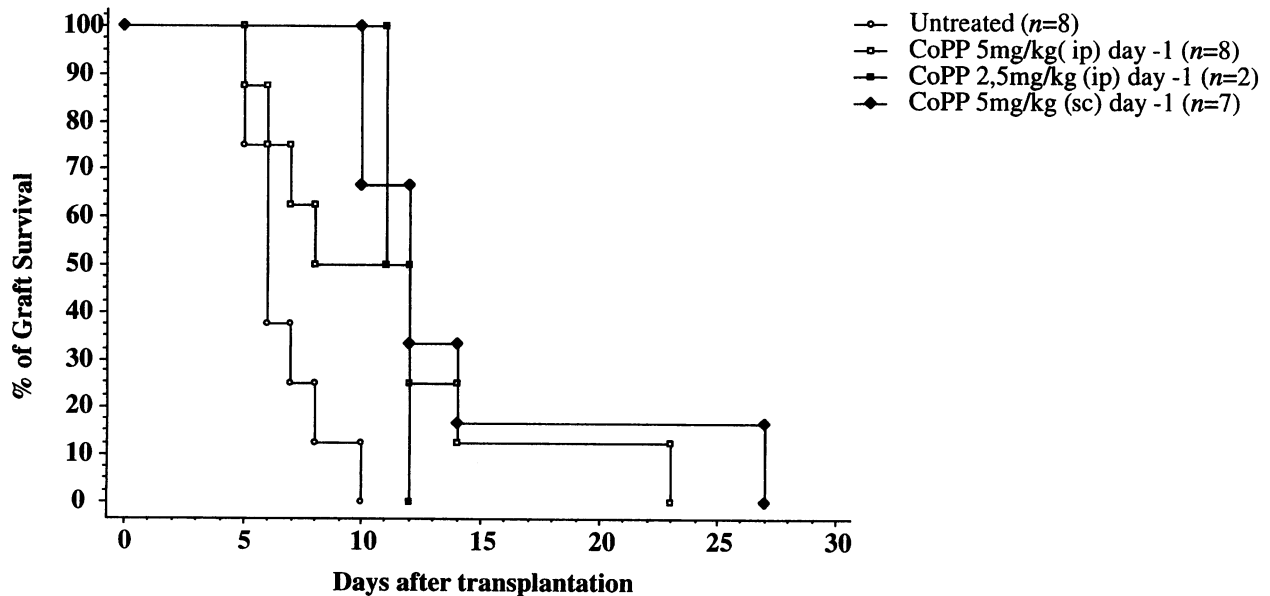
^c $p < 0.0008$ compared to untreated group, taking out the two >100 days from the analyses.

^d $p < 0.02$ compared to RDP1258 1 mg/kg IP day -7, -1 group.

Recipients were treated with either 5 or 2.5 mg/kg of CoPP (SC or IP) 1 day before transplantation. As shown in Figure 5, administration of CoPP resulted in a significantly prolonged heart allograft survival (mean allograft survival 13 ± 2.3 days vs. 6.6 ± 0.6 days; $p < 0.02$), though less than that observed with RDP1258.

Analysis of Graft-Infiltrating Cells

Graft-infiltrating cells (GIC) were analyzed 5 days after transplantation. We have previously shown that cellular infiltrate in LEW.1W heart allografts peaks on day 5 (14). Four animals in each group were therefore studied at this time



$p < 0.02$ untreated / CoPP 5mg/kg (sc) day -1

Fig. 5. Allograft survival after CoPP therapy. Kaplan-Meier depiction of heart allograft survival in rats receiving CoPP therapy.

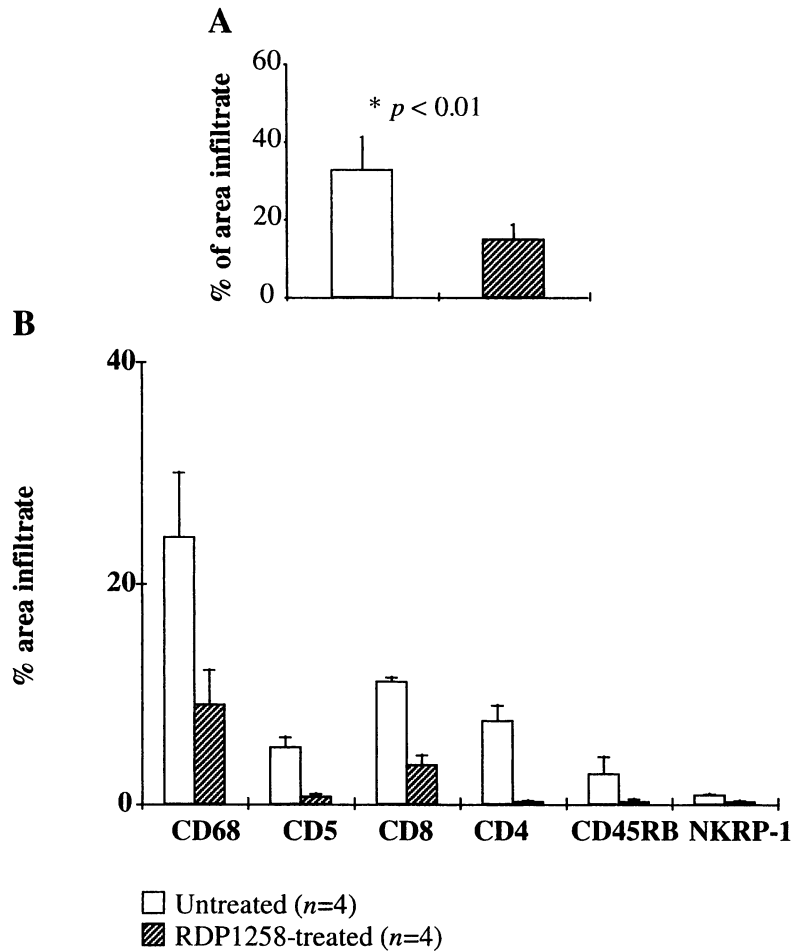


Fig. 6. Immunohistochemical analysis of graft infiltrating cells. (A) Sections from heart allografts removed from untreated or RDP1258-treated recipients 5 days after transplantation were stained with anti-CD45 specific monoclonal antibodies (OX1 and OX30). A quantitative analysis of graft infiltrating cells from untreated (open bar) and RDP1258 treated recipients (stippled bar) (mean \pm SD, $n = 4$) is shown; $*p < 0.01$. (B) Cryostat sections were stained with specific anti-rat monoclonal antibodies ED1 (CD68); OX19 (CD5, pan-T); W3/25 (CD4); OX8 (CD8); OX33 (CD45RB, B cells); 3.2.3 (NKRP-1) and quantified as described in Materials and Methods. The percentage of area infiltrated (mean \pm SD, $n = 4$) in untreated (open bars) and RDP1258-treated (stippled bars) is shown; $*p < 0.01$.

point; a quantitative analysis of cryostat sections is shown in Figure 6. A comparison of grafts from untreated and peptide treated recipients revealed a highly significant reduction in the number of GIC, labeled with OX1 and OX30 MAbs (anti-leukocyte common antigen CD45), ($32.7 \pm 9\%$ vs. $14.8 \pm 4\%$; $p < 0.01$) (Fig. 6A). As shown in Figure 6B, most GIC were CD68 positive (ED1 present in macrophages, monocytes, granulocytes, and dendritic cells), but T, B, and NK cells were also present. T cells were analyzed with OX19 (CD5, pan T lymphocytes), OX8 (CD8 $\alpha\beta$ and CD8 $\alpha\alpha$), and W3/25 (CD4); B cells with OX33 (CD45RA); and NK cells with MAb 3.2.3 (NKRP1). Numbers of all cell subsets studied were significantly reduced in grafts from peptide RDP1258-treated recipients (Fig. 6). The median of the percentage of area labeled with MAb CD68 was $24.2 \pm 6\%$ in untreated recipients vs. $8.9 \pm 3\%$ in RDP1258-treated recipients ($p < 0.01$); with MAb OX19, $5.2 \pm 1\%$ vs. $0.6 \pm 0.4\%$ ($p < 0.0002$); with MAb OX8, $11 \pm 0.6\%$ vs. $3.5 \pm$

1% ($p < 0.001$); with MAb W3/25, $7.5 \pm 1\%$ vs. $0.2 \pm 0.1\%$ ($p < 0.0001$); with MAb OX33, $2.7 \pm 1.6\%$ vs. $0.2 \pm 0.1\%$ ($p < 0.01$); and with MAb 3.2.3, $0.8 \pm 0.2\%$ vs. $0.1 \pm 0.2\%$ ($p < 0.01$).

Analysis of mRNA Expression in Allografts

To further assess the functional activity of GIC in allografts, we analyzed the expression of TNF- α , IFN- γ , IL-10, and iNOS. Messenger RNA levels were measured by competitive RT-PCR using RNA isolated from allografts harvested on day 5 after transplantation. Transcript numbers were standardized with respect to the expression of hypoxanthin phosphoribosyl transferase (HPRT) (Fig. 7). A comparison of iNOS mRNA levels in the grafts of four untreated and four peptide-treated recipients showed a significant increase in iNOS mRNA levels (0.25 ± 0.09 copies in untreated vs. 0.55 ± 0.09 copies in RDP1258 treated-recipients; $p < 0.05$). In contrast, TNF- α mRNA levels were significantly reduced follow-

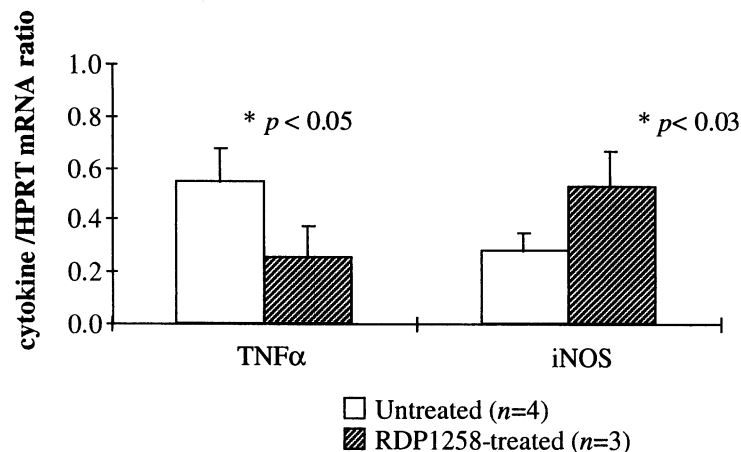


Fig. 7. Quantitative analysis of TNF- α and iNOS mRNA levels. Graft RNA was extracted from untreated (open bars) and RDP1258-treated (stippled bars) recipients 5 days after transplantation. RNA was reverse transcribed and analyzed by quantitative PCR. Results are expressed as mean \pm SD of the number of cytokine transcripts per number of HPRT transcripts.

ing peptide therapy (0.59 ± 0.05 copies in untreated vs. 0.28 ± 0.11 copies in RDP1258 treated-recipients; $p < 0.03$). Finally, analysis of IFN- γ (IFN- γ /HPRT ratio: untreated 0.027 ± 0.011 , $n = 4$; RDP1258-treated 0.042 ± 0.019 , $n = 4$) and IL-10, (IL-10/HPRT ratio: untreated 0.16 ± 0.05 , $n = 4$; RDP1258-treated 0.20 ± 0.09 , $n = 4$) revealed no significant differences in expression between the two experimental groups.

Cytotoxic Activity of GIC

We have previously (6) evaluated the peptide 07.75-84, derived from the $\alpha 1$ helix of HLA B7, in the LEW.1W/LEW.1A allograft model. Peptide 07.75-84 therapy prolonged heart allograft survival and reduced the cytotoxic activity of GIC (6). GIC isolated from RDP1258-treated recipients 5 days after transplantation were therefore tested both for specific anti-donor and nonspecific (NK cell-mediated) cytotoxicity in three independent experiments. In no instance was a reduction in GIC-mediated cytotoxicity observed following peptide RDP1258 therapy (Table 4).

Analysis of Alloantibody Production

Alloantibody production was analyzed on days 5 and 7 after transplantation. Antibody binding to donor type conA-stimulated donor cells expressing donor class I and II MHC antigens was measured using a serial dilution of decomplexed sera. IgG alloantibodies were evaluated with sera from four animals per group. No differences were observed in IgG antibody titers with sera from untreated or peptide-treated animals (Table 2).

Discussion

In this study, we evaluated the effect of a rationally designed immunomodulatory peptide on allograft rejection in a rat model. Peptide RDP1258 was shown to inhibit rat HO-1 in vitro in a dose-dependent manner. In vivo, administration of peptide RDP1258 resulted in up-regulation of splenic HO activity similar to that described with other known inhibitors of HO-1, in particular, cobalt-protoporphyrin. Peptide treatment of allograft recipients before transplantation resulted in significantly prolonged graft survival, a decrease in the number of graft infiltrating cells, decreased expression of graft TNF- α mRNA, and increased iNOS mRNA levels. No effects on IgG alloantibody responses were observed.

Both peptides 2702.75-84 and 07.75-84 derived from HLA class I have been shown to modulate HO-1 activity (10). Peptide 2702.75-84 in-

Table 4. Anti-donor and nonspecific cytotoxic activity of graft-infiltrating cells

	Donor cells (%)	YAC cells (%)
Untreated ($n = 3$)	22.8	15
RDP1258 treated ($n = 3$)	24	14

Graft-infiltrating cells from treated and untreated recipients were tested for alloantigen-specific (donor cells) and nonspecific (YAC cell line) cytotoxicity. Data shown are the mean of triplicate determinations of 25:1 ratio, spontaneous release being $<20\%$ of maximum release in all experiments.

hibits HO in the mouse, but not in the rat, whereas peptide 07.75-84 inhibits HO more effectively in the rat than in the mouse. This difference fits with the well-documented (8,16) specificity of the two peptides. Peptide 2702.75-84 has been shown to prolong allograft survival in mice but not in rats (7,25,26), whereas peptide 07.75-84 has been shown to prolong allograft survival in rats but not in mice (6,9,27,28). Peptide RDP1258, the design of which was based on *in vivo* data (prolonged graft survival), is a more effective inhibitor of both rat and mouse HO, and appears to prolong graft survival in both rodent species. CoPP, a well-known modulator of HO, has also been shown to prolong allograft survival in mice (24) and rats (this report). Taken together, these data support the hypothesis that modulation of HO activity is involved in peptide-mediated prolongation of allograft survival.

It is well established that inhibition of HO *in vivo* results in the up-regulation of HO mRNA and protein levels (19). This may be due to an increase in the concentration of intracellular heme, a toxic compound whose concentration has to be tightly controlled (19). Our peptide studies in mice are consistent with this observation (10). Administration of peptide into mice was shown to induce HO-1 gene transcription and HO-1 protein production. Depending on the half-life and potency of the administered HO-inhibitor, up-regulation of HO-1 in cells or animals is not necessarily associated with increased HO-1 activity. For example, administration of Zn-protoporphyrin, a very effective inhibitor of HO (21–23), resulted in up-regulation of HO-1 expression similar to the observations made with CoPP or peptide RDP1258. However, in contrast to CoPP and peptide RDP1258, injection of ZnPP resulted in a severe inhibition of HO activity *in vivo*. To further analyze the role of HO up-regulation in the *in vivo* effect of peptide, we treated the peptide-treated allograft recipients the day of transplantation with ZnPP in order to inhibit the HO activity. As shown in Table 3 ZnPP therapy completely inhibited the effect of the peptide therapy, which suggests a role for HO in peptide-mediated prolongation of allograft survival. Therefore, the up-regulation of HO-1 expression following administration of a HO inhibitor was expected to be dependent on the potency of the compound as well as its pharmacokinetics. This is consistent with our observations in rats. Peptide-mediated up-regulation of splenic HO expression and activity appeared to be dose dependent. The

highest increase in splenic HO activity was observed 1 day after administration of 1 mg/kg peptide. A fivefold increase in dose caused no further increase in splenic HO activity; instead, the initial increase in enzyme activity was lowered, while increased enzyme activity appeared to last longer. This may be due to the partial inhibition of HO following injection of a large dose of enzyme inhibitor (peptide RDP1258). This observation may explain why a high dose of peptide RDP1258 (5 mg/kg) was less effective in prolonging graft survival than a dose of 1 mg/kg when administered 1 day before transplantation and has a significant effect when administered 3 days before the graft (Table 3). In addition, it may explain why post-transplant therapy in our rat model was less effective than pretransplant therapy (data not shown). Furthermore, it could explain the differences observed with CoPP and peptide RDP1258. CoPP was shown to have a half-life of several days (24), whereas the half-life of peptide RDP1258 is expected to be short. Additional studies using various dose regimens will be necessary to clarify this issue.

Recently, up-regulation of HO-1 mRNA has also been described in renal allografts in rats 5 days after transplantation (29). Prominent expression of HO protein was observed in mononuclear cells infiltrating the renal allograft, >80% of the cells being macrophages. In addition to up-regulation of HO-1 transcription, an increased expression of the iNOS gene was observed. A close linkage between iNOS and HO expression has also been described in other models (19,30–32). These results are consistent with our observations in heart allograft recipients. Analysis of HO expression in spleen cells showed that the labeled cells were mostly macrophages and monocytes. Peptide therapy also caused increased HO expression in the spleen. Concomitantly, when iNOS expression in the graft was investigated, we found enhanced expression in peptide-treated recipients. In this context, it is interesting to note that a diet enriched in L-arginine, the substrate of nitric oxide synthase, improves graft survival in immunosuppressed rat allograft recipients (33). Furthermore, it has been suggested that up-regulation of iNOS expression confers partial protection on the aortic allograft from developing allograft arteriosclerosis (34).

Up-regulation of HO activity in mice has been shown to inhibit inflammation and several immune effector functions (24,35–39). In particular, up-regulation of HO-1 is associated with

decreased proliferation of lymphocytes and decreased production of TNF- α , INF- γ , and IL-10. In contrast, production of IL-2 was not affected. Inhibition of HO-1, on the other hand, has been shown to stimulate the production of TNF- α and INF- γ (24,39). Preliminary results of experiments with purified macrophages/monocytes and T cells from CoPP-treated animals indicated that the modulation of T cell effector functions is dependent on the presence of macrophages/monocytes (J. Woo, unpublished observations). These observations are consistent with our finding that peptide treatment resulted in a decreased expression of TNF- α in the graft.

Peptide therapy caused a reduction in the number of graft-infiltrating cells analyzed on day 5 post-transplantation. While it is unclear why peptide administration influenced graft infiltration, several possibilities can be envisioned. (i) Peptide-mediated immunosuppression may result in decreased production of cytokines and chemokines by graft-infiltrating lymphocytes. Consequently, less cells are routed into the allograft. (ii) Reduced cytokine production may result in decreased activation of endothelial cells. (iii) Peptide therapy may not only modulate immune effector functions but also directly activate endothelial cells. Further studies will be necessary to clarify these questions.

Peptide RDP1258 was shown to inhibit cytotoxic T and NK cells more effectively than peptide 07.75-84. We have previously reported a reduction in T and NK cell-mediated cytotoxicity following peptide 07.75-84 treatment of recipients (6,16). However, analysis of GIC showed no difference in cytotoxic activity following peptide RDP1258 therapy. This may be due to differences in pharmacokinetics and dose responses. As outlined above, such differences could have profound effects on modulation of HO activity in vivo. On the other hand, we cannot exclude the possibility that peptides 07.75-84 and RDP1258 do not have identical biological activity and further studies will be necessary to clarify this issue.

Treatment with CoPP has also been shown to lower cell-mediated cytotoxicity (24). However, this effect required the administration of a large dose of CoPP (20 mg/kg) on 3 consecutive days, suggesting that it may be due not to up-regulation of HO activity but only to the modulation of other enzymes. Indeed, in addition to its effect on HO, Cobalt-protoporphyrin has been shown to modulate other heme enzymes, such as NOS and guanylate cyclase (40,41). Therefore, high doses of CoPP may modulate several of the bio-

chemical pathways influencing immune effector functions. While we cannot exclude the possibility that peptide RDP1258 may influence various different enzymes and pathways as well, our results support the hypothesis that its mechanism of action involves modulation of HO activity. In particular, up-regulation of HO-1, inhibition of the proliferation of lymphocytes, and reduced production of inflammatory cytokines may all be closely linked. How HO-1 up-regulation causes these effects remains unclear. However, all of the endproducts of heme degradation—biliverdin, bilirubin, and carbon monoxide—have been shown to modulate immune effector functions. Biliverdin has been shown to inhibit human complement (42) phytohemagglutinin-induced proliferation, IL-2 production, and antibody-dependent and -independent cell-mediated cytotoxicity (43,44). Carbon monoxide, like NO, has been shown to stimulate the production of cGMP (45,46). The secondary messenger, cGMP, is involved in cell growth arrest and TNF- α release by macrophages (47). In addition, cGMP is involved in the regulation of various protein kinases, phosphodiesterases, and ion channels (48–50). Carbon monoxide has also been shown to inhibit NO synthase models (18,30–32). On the basis of these observations, multiple pathways of immunomodulation involving HO may be envisioned, the further characterization of which may lead to the development of novel strategies for modulation of the immune system.

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