# Pattern of Hypocretin (Orexin) Soma and Axon Loss, and Gliosis, in Human Narcolepsy

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Human narcolepsy is correlated with a greatly reduced number of hypocretin (orexin) containing neurons and axons, and an elevated level of hypothalamic gliosis. We now report that the percentage loss of Hcrt cells and percentage elevation of GFAP staining are variable across forebrain and brainstem nuclei, and are maximal in the posterior and tuberomammillary hypothalamic region. Regional gliosis and percent loss of hypocretin axons in narcoleptics are not correlated with regional hypocretin cell soma density in normals or with regional percent soma loss in narcoleptics. Rather they are independently and strongly correlated with the regional density of hypocretin axons and the message density for hypocretin receptor 2, as guantified in the rat. These results are consistent with the hypotheses that the loss of hypocretin function in narcolepsy results from a cytotoxic or immunologically mediated attack focused on hypocretin receptor 2 or an antigen anatomically linked to hypocretin receptor 2, and that this process is intensified in regions of high axonal density.

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#### Introduction

We previously reported that human narcolepsy patients had an 85 to 95% reduction in the number of hypocretin (Hcrt) neurons (45, 46). At the same time, Peyron et al (39) reported that the level of preprohypocretin mRNA in narcoleptics was below their detection threshold. In our prior work, we also noted the presence of gliosis in the Hcrt cell region.

The cause of Hcrt cell loss in narcolepsy remains unclear. Narcolepsy is associated with the presence of a particular human leukocyte antigen, DQB1\*0602, with more than 90% of narcoleptics having this haplotype, compared to 20 to 30% of the overall population (12, 15, 26, 31, 33). This association and the presence of gliosis within Hcrt cell regions of narcoleptics (46) suggests that Hcrt cell loss may be caused by an inflammatory or immune mediated process rather than by a developmental abnormality (11, 43). A better understanding of the pattern of damage to Hcrt containing somas and axons and of the gliosis occurring in narcolepsy should clarify the nature of the pathological process responsible for this disorder.

In the current study we have quantified the anatomical distribution of Hcrt cell loss, the reduction in numbers of Hcrt containing axons and the increased numbers of astrocytes in narcoleptics. We have examined the correlations between these measures and also compared the extent of cell loss and gliosis to the reported density of the 2 known Hcrt receptor mRNA's, quantified in the rat (30, 49). These data have been reported in preliminary form (48).

#### **Materials and Methods**

Narcolepsy was diagnosed according to standard criteria (10). In the current study we quantified the histological changes in 5 narcoleptic patients, including 4 used in our prior study that found Hcrt cell loss and gliosis in narcolepsy, and 6 matched normals. We did not receive brain stem tissue from patient ND (Table 1). Immunohistochemistry and antigen retrieval for Hcrt staining on the brains that had been in fixative for an extended period of time was done as previously described (19, 46). The hypothalamic and brainstem tissue blocks were cut into 40-µm thick coronal sections with a microtome. The sections were treated with 0.5%sodium borohydride in PBS for 30 minutes and washed with PBS, and then incubated for 30 minutes in 0.5% H<sub>2</sub>O<sub>2</sub> for blocking of endogenous peroxidase activity. For antigen retrieval, sections were heated for 30 minutes at 80°C in a water bath with 10 mM sodium citrate (pH 8.5) solution. The sections were cooled to room temperature in sodium citrate and washed with PBS. Antigen retrieval was not required on narcoleptic brain NE, which had been in fixative for only 3 months or for GFAP staining in any of the brains. After thorough washing with PBS, the sections were placed for 2 hours in 1.5% normal goat serum in PBS and incubated for 72 hours at 4°C with a 1:2000 dilution of rabbit anti-Hcrt-

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Subject	Sex	Age	Cause of death	Time in Fix years	Disease Onset age	Medications	Cataplexy
Narcoleptics							
NA	F	63	adenocarcinoma	5.0	20	Clomipramine, Vivactil	+
NB	Μ	49	cardiovascular	4.5	18	Dexedrine, Vivactil, GHB	+
NC	Μ	60	sepsis	4.2	23	Ritalin, Dexedrine	+
ND	М	86	pulmonary	12.0	35	Ritalin	-
NE	М	81	cardiovascular	0.3	13	Dexedrine	+
Normals							
CA	F	54	liver failure	0.5			
CB	F	42	AML/DIC	1.0			
CC	Μ	n/a	non-neurological	1.0			
CD	Μ	73	pulmonary, cv	5.0			
CE	Μ	n/a	non-neurological	16.0			
CF	Μ	n/a	non-neurological	16.0			

cv, cardiovascular; AML, acute myelocytic leukemia; DIC, disseminated intravascular coagulation; GHB, gamma-hydroxy butyrate; "non-neurological" indicates no history of neurological illness and no specified cause of death.

Table 1. Characteristics of narcoleptic and normal subjects.

1 or rabbit anti-Hcrt-2 (Orexin A and B, Oncogene Research Products). Sections were then incubated in a secondary antibody (biotinylated goat anti-rabbit IgG; Vector Laboratories) followed by avidin-biotin peroxidase (ABC Elite Kit; Vector laboratories), for 2 hours each at room temperature. The tissue-bound peroxidase was visualized by a diaminobenzidine reaction (Vector laboratories). We used a 15-minute diaminobenzidine incubation on all tissues, having determined in pilot studies on normal and narcoleptic human tissue that this duration produced optimal staining with cell boundaries clearly visible and minimal background artifact. Narcoleptic and normal tissue were always stained in pairs along with a no-antibody control. For identification of activated astrocytes, glial fibrillary acidic protein (GFAP) was immunostained with a 1:2000 dilution of primary polyclonal rabbit anti-cow GFAP antibody (DAKO, Carpinteria, Calif). Counterstaining of adjacent sections with cresyl violet aided in the localization of immunoreactive elements in relation to nuclei.

Brain regions and nuclei were identified using the atlas *Structure of the Human Brain* by DeArmond et al, 1989 (7). The following nuclei were examined: anterior hypothalamus (AH), periventricular nucleus (PVN), paraventricular nucleus (PAVN), supraoptic (SO), arcuate nucleus (ARN), dorsal hypothalamus (DH), dorso-medial hypothalamus (DMH), ventromedial hypothalamus (VMH), lateral hypothalamus (LH), posterior hypothalamus (PH), tuberomammillary nucleus (TMN), mammillary nucleus (MN), thalamus (dorso-medial nucleus, TH), locus coeruleus (LC), raphe dorsalis (RAD) and raphe centralis (RAC). Hcrt labeled

Hort cell density in hypothalamus



**Figure 1.** Hypocretin (Hcrt) soma density is greatly reduced in human narcolepsy, with maximal percent loss in posterior hypothalamus. Density of Hcrt cells in the hypothalamus of normal and narcoleptic brains: anterior hypothalamus (AH), dorsal hypothalamus (DH), dorsomedial hypothalamus (DMH), posterior hypothalamus (PH), lateral hypothalamus (LH). (\*\* p<0.01, \*\*\*\*p<0.0001, \*\*\*\*\*p<0.00001, Student's *t* test, df=6).

cells and axons, and GFAP labeled cells were counted "blind" on a Neurolucida computerized microscope interface. We used a Nikon E600 microscope with three axis motorized stage, video camera, and Stereoinvestigator software (MicroBrightfield Corp., Colchester, Vt). Digital image acquisition was carried out with a Spot camera (Diagnostic Instruments, Sterling Heights, Mich) and imported to the Corel Draw program. Contrast and brightness were corrected. Hcrt soma and axon loss and observed gliosis were compared with each other and with the relative density of Hcrt receptor mRNA expression reported for the rat (30).

Data analysis. Hypothalamic tissue from all 5 narcoleptic brains (NA-NE) and 6 matched controls (CA-CF) was used for quantification of Hcrt cells, axon and GFAP stained astrocytes. Brainstem tissue from 4 narcoleptic (NA-NC and NE) and 4 neurologically normal individuals (CC-CF) was used for Hcrt and GFAP staining. To obtain unbiased number and size estimates from a sample of axons in each nucleus, axon profiles of any shape, size and orientation and in all locations must have an equal probability of being sampled. The unbiased counting frame provides rules for defining which profiles belong to the area of the counting frame. In a complete tessellation of unbiased counting frames, all axon profiles were sampled once and only once. Systematic, uniform, random sampling with counting frames across the entire nucleus ensures that all locations within the nucleus are equally represented and that all axon profiles are sampled with equal probability. Hcrt and GFAP cell number were determined with stereological techniques on a one in 6 series of 40-µm sections through the Hcrt containing regions of the hypothalamus (16, 46) and brain stem nuclei. The numerical densities of the cells were calculated as number of cells per unit area (mm<sup>2</sup>). The number of axons counted with the unbiased counting frames yields an unbiased estimate of the total number of axons (25). We used  $250 \times 250 \ \mu m$ as the counting frame size. By identifying the axon hillock or by excluding tapering processes each axon could be distinguished from dendritic branches in the regions containing Hcrt soma. Each axon and its branches was counted as a single item. The numerical density of Hcrt axons were calculated as number of axons per unit area (mm<sup>2</sup>). All values for the each nucleus were calculated for each subject. These were pooled to give means and SD for each region and each group. Once the numerical densities were obtained, the density and percentage change of axons, somas and GFAP labeled cells was calculated. The data were analyzed using a one-way analysis of variance (ANOVA). All "t" values reported are Bonferroni adjusted. Statistical significance was set at P<0.05.

*Partial correlation analysis.* In order to statistically separate the effects of cell loss, fiber loss, Hert receptor density and gliosis, we performed a series of partial correlation analyses. The relative message density for receptor 1 or 2 was taken from Marcus et al (30). The message density was rated on a scale from 0 to 3: 0 applied to areas nearly entirely devoid of receptor; 1 applied to areas with low density; 2 applied to areas with moderate density; and 3 applied to areas with

highest density. For the case of the partial correlation of Z (eg, message density for type 2 receptor) on Y (eg, % of fiber loss) controlling X (eg, gliosis), the bivariate regression of Z on X was performed, so that the residuals were the variance in Z after the effect of X was considered. The bivariate regression of Y on X was performed, so that the residuals were the variance in Y after the effect of X is considered.

### Results

All 5 narcoleptic brains, including one in which antigen retrieval had not been performed, showed the same pattern of Hcrt cell and axonal loss, and gliosis. To control for the time spent in fix, we processed 3 neurologically normal brains that, like the narcoleptic brains, had been held in fix in our laboratory for 5 or more years and stained them with the same antibodies. As in the narcoleptic brains, no Hcrt staining was apparent in these normal brains unless our antigen retrieval procedure was followed. All normal brains, including the three in which antigen retrieval had been performed showed the same pattern of Hcrt cell and axon staining.

Hert cell loss in the narcoleptic hypothalamus. Hert immunoreactive cells were found in anterior (AH), dorsal (DH), dorsomedial (DMH), posterior (PH) and lateral (LH) hypothalamic nuclei of both normal and narcoleptic brains as previously reported (32, 45, 46) (Figure 1). Cell number was significantly reduced in narcoleptics compared to normals (p<0.00001, F=238.35, df=1, 30) and there was a significant interaction between hypothalamic nucleus and the percentage of Hert cells lost (p < 0.00001, F = 33.1, df = 4, 30), indicating that percentage cell loss was significantly more severe in certain nuclei. There was no significant correlation between Hcrt cell density in normals and percent of Hert cell loss in narcoleptics (p<0.15, r=0.54, t=1.91, n = 10). In the normal brain, Hert cell density was highest in DMH (41 cells/mm<sup>2</sup>) and lowest in AH (2 cells/mm<sup>2</sup>). In narcoleptics, the maximum percentage loss of Hcrt cells occurred in the posterior hypothalamus (97%, p<0.0004, t=7.9, df=6; Figure 2). The minimum percent loss of Hcrt cells was seen in AH (74%, p<0.022, t=3.6, df=6). Overall, narcoleptics had a mean 89% reduction in Hcrt cell number (range 74-97% between subjects) compared to the average number seen in normals.

*Hcrt axon loss in the narcoleptic brain.* Hcrt immunoreactive axons were found throughout the hypothalamic and brainstem nuclei of both normal and

### NORMAL

## NARCOLEPTIC



Figure 2. Hcrt cells in the hypothalamic nuclei of normal and narcoleptic humans. The abbreviations are same as in Figure 1. Scale  $bar = 50 \mu m$ .

narcoleptic brains (Figures 3A, 4; Table 2). In the hypothalamus of normal brains, a high density of Hcrt axons was found in paraventricular (69.7 axons/mm<sup>2</sup>), periventricular (59.0 axons/mm<sup>2</sup>), posterior (48.7 axons/mm<sup>2</sup>), dorsomedial (48.5 axons/mm<sup>2</sup>) and tuberomammillary nuclei (47.5 axons/mm<sup>2</sup>). A low axon den-



**Figure 3.** Density of Hcrt axons and GFAP labeled cells varies across nuclei between normal and narcoleptic brains. **A.** Hcrt axon density (axons/mm<sup>2</sup>) in normal and narcoleptic brain arranged according to descending magnitude in normal brain. **B.** GFAP density (cells/mm<sup>2</sup>) in normal and narcoleptic brain arranged according to descending magnitude in narcoleptic brain. (\*p<0.05, \*\*p<0.01, \*\*\*p<0.001, \*\*\*\*p<0.0001, \*\*\*\*p<0.0001, student's ttest, df = 6). Paraventricular nucleus (PAVN), periventricular nucleus (PVN), locus coeruleus (LC), posterior hypothalamus (PH), dorsomedial hypothalamus (DMH), tuberomammillary nucleus (TMN), raphe dorsalis (RAD), ventromedial hypothalamus (VMH), anterior hypothalamus (AH), dorsal hypothalamus (DH), arcuate nucleus (ARN), raphe centralis (RAC), lateral hypothalamus (LH), supraoptic (SO), mammillary nucleus (MN), thalamus (TH).

Nucleus	Axon Density	% Loss in	Hcrtr-2	GF/	AP Name	GFAP
	Normais (axons/mm²)	Narcoleptics	density	Narcoleptics (cells/	mm²)	% Increase
AH	39.0 ± 1.7	65.02	1	39.0 ± 5.7	19.6 ± 2.2	98.38
PVN	$59.0 \pm 5.6$	71.69	2	$43.3 \pm 9.7$	21.3 ± 3.1	103.05
PAVN	$69.6 \pm 6.7$	76.89	3	59.9 ± 10.3	17.7 ± 4.8	238.61
SO	$20.7 \pm 6.6$	54.12	1	$12.4 \pm 4.6$	7.8 ± 1.2	57.84
ARN	35.7 ± 8.1	69.49	3	$37.3 \pm 5.8$	14.9 ± 1.2	150.10
ЭΗ	$36.8 \pm 2.2$	67.99	1	40.5 ± 14.1	14.7 ± 2.6	174.50
ОМН	$48.4 \pm 9.8$	72.53	2	34.6 ± 15.5	18.3 ± 3.5	89.05
/MH	39.1 ± 9.5	72.58	2	36.8 ± 11.8	$16.2 \pm 3.0$	127.29
H	$30.5 \pm 8.2$	71.38	3	$35.8 \pm 7.3$	13.6 ± 2.1	161.70
РΗ	$48.7 \pm 6.4$	74.04	2	46.3 ± 15.1	$10.5 \pm 3.3$	340.74
ΓMN	$47.4 \pm 6.4$	71.44	3	41.7 ± 7.3	14.7 ± 5.3	183.54
MN	$6.9 \pm 3.5$	52.46	1	8.1 ± 2.4	$7.2 \pm 2.9$	12.96
ГН	$2.5 \pm 0.6$	22.20	0	7.9 ± 1.2	$7.5 \pm 2.2$	5.61
RAC	$35.3 \pm 7.1$	49.64	0	45.8 ± 5.5	38.12 ± 3.0	20.21
RAD	$45.9 \pm 7.0$	53.15	1	60.1 ± 3.0	50.48 ± 8.7	19.24
_C	$53.2 \pm 6.2$	64.32	0	61.2 ± 9.1	30.21 ± 9.5	102.71
Vean	38.70 ± 2.7	63.06	1.56	38.20 ± 4.4	18.95 ± 2.3	117.86

Table 2. Axon density in normals, percent loss in narcoleptics, message density for Hcrt receptor (Marcus et al, 2001) and GFAP percent increase in narcoleptics in surveyed nuclei.

sity was observed in mammillary nucleus (6.9 axons/mm<sup>2</sup>) and thalamus (dorsomedial nucleus, 2.5 axons/mm<sup>2</sup>). In the brainstem, locus coeruleus had the highest density of Hcrt axons (53.3 axons/mm<sup>2</sup>). Raphe centralis and raphe dorsalis had 46.0 and 35.4 axons/mm<sup>2</sup> respectively. Compared to normals, nar-

coleptics had a significant reduction in the number of Hert axons in brainstem nuclei and hypothalamic nuclei (p<0.00001, F=720.42, df=1, 96) and there was a significant interaction between nucleus and percent axon loss (p<0.00001, F=12.02, df=15, 96), indicating that percent axonal loss was significantly greater in certain

**Figure 4.** (Opposing page) Hcrt axons in hypothalamus and brainstem nuclei of normal and narcoleptic brains: paraventricular (PAVN), periventricular (PVN), locus coeruleus (LC), tuberomammillary (TMN), raphe dorsalis (RAD) ventromedial hypothalamus (VMH), raphe centralis (RAC), arcuate nucleus (ARN), supraoptic (SO), mammillary (MN), thalamus (TH). Scale bar=50 μm.



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**Figure 6.** Pattern of gliosis and percentage of Hcrt axon loss in the hypothalamus and brain stem nuclei of normal and narcoleptic brain. Number of GFAP labeled cells is correlated with percent Hcrt axon loss. The abbreviations are same as in Figure 3.

nuclei. Among the hypothalamic nuclei, PAVN (77%, p < 0.00044, t = 14.68, df = 6), PH (74%, p < 0.0023, t =7.3, df=6), VMH (73%, p<0.0075, t=5.86, df=6), DMH (73%, p<0.0031, t=6.9, df=6), TMN (71%, p<0.00059, t=9.3, df=6), PVN (72%, p<0.00029, t= 10.57, df=6) and LH (71%, p<0.014, t=5.19, df=6) had a high percentage reduction of Hcrt axons (Figure 3A). There was a significant, but lesser percent reduction of Hcrt axons in LC (64%, p<0.0014, t=6.89, df=6), RAD (53%, p<0.008, t=4.8, df=6) and RAC (50%, p < 0.014, t = 4.36, df = 6). The percent axon density reduction in Hert innervated nuclei in narcoleptics was positively correlated with the density of Hcrt axons in normal humans (r=0.75, t=3.37, p<0.0008, n=11). The total number of Hcrt axons in all the structures analyzed was reduced by 67% compared to an 89% reduction of Hert soma count in the same brains, suggesting that either Hcrt cells with smaller axonal fields are lost to a greater extent in narcolepsy or that some sprouting of axons of surviving Hcrt cells may have occurred in narcoleptics.

*GFAP labeled astrocytes.* GFAP labeled astrocytes were found throughout the hypothalamic and brainstem nuclei of normal and narcoleptic brains (Figures 3B, 5). In the normal brain, compared with hypothalamic nuclei, locus coeruleus, raphe dorsalis and centralis had a higher density of GFAP stained cells (Figure 3B; Table 2). There was a significant increase in gliosis indicated by GFAP staining (p<0.00001, F=243.86, df



**Figure 7.** The pattern of gliosis and Hcrt axon loss ranked in relation to the message density for Hcrt receptors. **A.** The percent increase in gliosis and axon loss was not correlated with the message density for Hcrt receptor 1. **B.** The percent increase in gliosis and percent axon loss was correlated with the message density for Hcrt receptor 2. The abbreviations are same as in Figure 3. The message density for receptor data from previously reported data (Marcus et al, 2001, +++ highest density, ++ moderate density, + low density, - absent).

= 1, 102) and a significant interaction between nucleus and amount of gliosis in narcoleptics (p<0.00001, F=5.65, df=15, 102) indicating that the increase in GFAP staining was significantly greater in certain nuclei. The size of GFAP labeled astrocytes did not differ between narcoleptics and controls. In narcoleptics, GFAP density was high in paraventricular (60.0 cells/mm<sup>2</sup>), posterior (46.3 cells/mm<sup>2</sup>), periventricular (43.3 cells/mm<sup>2</sup>), tuberomammillary (41.8 cells/mm<sup>2</sup>), and dorsal nuclei of hypothalamus (40.5 cells/mm<sup>2</sup>). Compared to normals,

Figure 5. (Opposing page) GFAP labeled astrocytes in the hypothalamus and brainstem nuclei of normal and narcoleptic brain. The abbreviations are same as in Figure 3. Scale bar = 50  $\mu$ m.

Y = % axon   Z = message	oss narcoleptic e density for Hci	s t type 2 recepto	r					
Simple rxy rxz ryz	r 0.752 0.426 0.724	r <sup>2</sup> 0.565 0.181 0.523	<b>P</b> 0.0008 0.0995 0.0015	Partials rxy.z rxz.y ryz.x	r 0.682 -0.181 0.644	r <sup>2</sup> 0.466 0.033 0.415	t 3.37 -0.66 3.04	<b>P</b> 0.0050 0.5207 0.0094
Conclusions 1. Hcrt axon	: loss is correlate	ed with normal fi	ber density		_			
2. Hcrt axon 3. These cor	loss is correlate relations are inc	ed with the mess dependent	sage density for H	icrt type 2 recepto	r			
2. Hcrt axon 3. These cor <b>B</b> X = gliosis (0 Y = axon dei Z = message	loss is correlate relations are inc GFAP) percent i nsity normals e density for Hor	ad with the mess dependent ncrease in narco t type 2 recepto	sage density for F		r			
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2. Hcrt axon 3. These cor <b>B</b> ( = gliosis (( 7 = axon dei 2 = message <b>Simple</b> rxy rxz	GFAP) percent in asity normals density for Hor r 0.559 0.632	ad with the mess dependent ncrease in narco t type 2 recepto r <sup>2</sup> 0.312 0.399	Deptics 0.0244 0.0086	Partials rxy.z rxz.y	r 0.413 0.525	r² 0.171 0.276	t 1.64 2.22	<b>P</b> 0.1249 0.0448
2. Hcrt axon 3. These cor <b>B</b> <b>C</b> = gliosis (C <i>C</i> = axon der <i>C</i> = message <b>Simple</b> rxy rxz ryz	GFAP) percent i nsity normals e density for Hcr 0.559 0.632 0.426	ncrease in narco t type 2 recepto r <sup>2</sup> 0.312 0.399 0.181	Deptics r 0.0244 0.0086 0.0995	Partials rxy.z rxz.y ryz.x	r 0.413 0.525 0.113	r <sup>2</sup> 0.171 0.276 0.013	t 1.64 2.22 0.41	<b>P</b> 0.1245 0.0448 0.6884
2. Hcrt axon 3. These cor <b>B</b> K = gliosis (C Y = axon der Z = message <b>Simple</b> rxy rxz ryz Conclusions	GFAP) percent i nsity normals e density for Hor 0.559 0.632 0.426	ncrease in narco t type 2 recepto r <sup>2</sup> 0.312 0.399 0.181	Deptics r 0.0244 0.0086 0.0995	Partials rxy.z rxz.y ryz.x	r 0.413 0.525 0.113	r <sup>2</sup> 0.171 0.276 0.013	t 1.64 2.22 0.41	<b>P</b> 0.124§ 0.044§ 0.6884

Table 3. Partial correlation analysis of axonal loss, axonal density and GFAP staining.

narcoleptics had a very high percentage increase of GFAP in posterior hypothalamus (341%, p<0.017, t= 4.65, df=6), paraventricular (239%, p<0.025, t=4.28, df =6), tuberomammillary (184%, p<0.0049, t=5.9, df=6), dorsal hypothalamus (175%, p<0.056, t=3.56, df=6), and arcuate nucleus (150%, p<0.0014, t=7.4, df=6). In supraoptic, mammillary and thalamic nuclei we found a very low density of GFAP in both normal and narcoleptic brains (Figure 3B). Locus coeruleus had a higher percentage of GFAP increase (103%, p<0.0026, t= 5.14, df = 8) than raphe dorsalis (32%, p < 0.59, t = 1.4, df=8) and raphe centralis (20%, p<0.061, t=2.8, df=8)(Figure 6), and brainstem nuclei in general showed smaller increases than forebrain nuclei. Regional gliosis in narcoleptics was highly correlated with the density of Hert axons in the normal brain (r=0.87, p<0.0001, n=11).

Hcrt soma and axon loss and message density for Hcrt receptors. The reported anatomical distribution of Hcrt receptor 1 and 2 in the rat are largely distinct (Figure 7 A, B). Hcrt soma loss was not correlated with the message density reported for Hcrt receptor 1 (p<0.66, r =0.26, t=0.47, n=10) or with that reported for Hcrt receptor 2, as quantified in the rat (30) (p<0.53, r=0.37, t=0.69, n=10), across the hypothalamic nuclei. The percentage of Hcrt axon loss in narcoleptics was compared with message density for Hcrt receptors. Hcrt axon loss was positively correlated with the message density for Hcrt receptor 2 (r=0.72, t=10.95, p<0.002). Axon loss was not correlated with the message density for Hcrt receptor 1 (r=0.05, t=10.55, p<0.84).

**Partial correlation analysis.** In order to statistically separate the effects of cell loss, fiber loss, message density for receptor and gliosis, we performed a series of partial correlation analyses (Table 3A, B). These analyses showed that Hcrt axonal density in normals correlated positively with percent axonal loss in narcoleptics over all examined nuclei. They also showed that the message density for Hcrt receptor 2, as measured in the rodent, correlated positively with the percent axonal loss in human narcoleptics. These 2 correlations were found to be independent. The increase in gliosis in narcoleptics was also shown to be correlated independent-ly with message density for type 2 receptor independent of normal axon density. In summary, we identify 2 independent correlates of axonal loss and gliosis in nar-

coleptics, the density of axonal fibers and the message density for Hcrt type 2 receptor.

### Discussion

We find the highest percentage depletion of Hcrt cells and GFAP staining in the posterior and tuberomammillary hypothalamic regions. This region contains all of the hypothalamic histamine cells (1, 27, 37, 44). Hypocretin potently excites histamine neurons (3, 9,54). A reduction of histamine has been seen in the cerebrospinal fluid of genetically narcoleptic dogs (35) and narcoleptic humans (36). It is well known that antihistamines cause sleepiness and that stimulation of the histamine cells produces arousal (28). The arousing effect of Hert has been shown to be heavily dependent on activation of histamine containing cells (18). The loss of Hcrt cells in human narcoleptics suggests that reduction in Hcrt activation of histamine cells may mediate some of the symptoms of sleepiness in narcolepsy. The increased percentage loss of Hcrt cells, axons and gliosis in the posterior hypothalamus, suggests that this region may be a focus of inflammation and damage in narcolepsy.

In contrast to our prior report (46) and current findings, a prior study (Peyron et al, 2000) did not detect gliosis in the hypothalamus of the 2 narcoleptic brains inspected or in any control tissue. The relative sensitivities of the staining techniques and procedures used may account for this difference between the Peyron et al study and the current work.

The distribution of Hcrt cell somas and axonal projections is quite similar in mice (8, 41), rats (34, 40), cats (52), dogs (47) and humans (46). In the current study, we correlated the levels of cell and axonal loss with prior published work on Hcrt mRNA receptor 1 and 2 message density in the rat (30, 49). Comparable receptor data are not available for humans. In general, receptor type distributions, like soma and axonal distributions, are preserved across species (13, 14, 17, 20, 42, 51), so it is likely that the reported Hcrt receptor data in the rat approximate that in the human. The very high correlations seen here would be less likely if the rodent receptor density was not linked to a meaningful neurological parameter in humans, most likely receptor density. However, we cannot rule out the possibility that the underlying determinant of the relationships we see here is a function of some other correlate of rodent Hcrt receptor 2 mRNA concentration.

Our partial correlation analyses lead to the following conclusions: *i*) percentage Hcrt axon loss/increase in gliosis is not correlated with the density of Hcrt cell somas; *ii*) percentage Hcrt axon loss is correlated with

Hert axon density; *iii*) percentage Hert axon loss is also correlated with the message density for Hert receptor 2; *iv*) percentage increase in gliosis is correlated with the message density for Hert receptor 2; and *v*) these correlations are independent.

The correlation between percentage loss of axons and percentage increase in gliosis, and the message density for Hcrt receptor 2 suggests that this receptor or antigens linked to it may be associated with processes that intensify the pathological process in narcolepsy.

Canine genetic narcolepsy is caused by one of 2 distinct mutations of the Hcrt receptor 2, but not by mutations of Hert receptor 1 (29). Similarly, knockouts of the Hcrt 2 receptor, but not the Hcrt 1 receptor produce symptoms of narcolepsy in mice (6, 21). Our current finding that damage in human narcolepsy is correlated with the best available estimate of Hcrt receptor 2 message density suggests that damage in the region of this receptor is a determining factor in human narcolepsy. Both cell loss in humans (46) and animals (11) and receptor mutation (29, 39) lead to the similar syndromes of excessive sleepiness during the active period of the circadian cycle and disrupted sleep during the inactive phase, cataplexy, and, as studied in dogs and humans, therapeutic response to anticholinergic drugs and to drugs that enhance monoamine levels, and symptom exacerbation by cholinesterase inhibitors and the alpha-1 antagonist prazosin (2, 5, 43, 53). The striking similarity of the symptoms and pharmacology of narcolepsy in Hert receptor 2 mutant animals to those in hypocretin deficient humans (46) suggests that normal receptor 2 operation is necessary for adequate regulation of Hcrt release at terminal sites.

In genetically narcoleptic dogs, we have seen an abrupt termination of activity in locus coeruleus in conjunction with cataplexy (53). In normal animals, locus coeruleus cells never have a comparable cessation of activity in waking, and have an important role in muscle tone facilitation (24). Although the locus coeruleus has the densest extrahypothalamic concentration of Hert axons (40), the predominant receptor type in this region is Hcrt receptor 1 (30, 49). Therefore the Hcrt receptor 2 mutation effect in genetically narcoleptic dogs must be indirect. Thus, we hypothesize that the Hcrt receptor 2 dysfunction disrupts the normal pattern of Hcrt release into locus coeruleus and other sites during waking. In human narcolepsy the loss of Hcrt projections to the locus coeruleus would have a similar effect. We have found that Hcrt -1 release in hypothalamic regions is high during REM sleep, equally high in waking, but significantly reduced in nonREM sleep (22, 23). It is likely that Hcrt release interacts with GABA and glutamate to produce differing state specific postsynaptic effects (4, 38, 50) thereby modulating arousal and motor control.

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