Postglucose growth hormone nadir and insulin-like growth factor-1 in naïve-active acromegalic patients: do these parameters always correlate?

Nadir do hormônio de crescimento após glicose e fator de crescimento insulina-símile do tipo 1 em pacientes acromegálicos ativos virgens de tratamento: são estes parâmetros sempre correlacionáveis?

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The diagnosis of acromegaly is based on clinical features and biochemical criteria. While clinical suspicion is of extreme importance, laboratory evaluation is mandatory to establish the diagnosis. Growth hormone (GH) is secreted in a pulsatile pattern. Normal individuals have undetectable levels during most of the daytime period. In contrast, GH is always detectable in acromegalic patients, promoting higher levels of insulin-like growth factor-1 (IGF-1) (1). Thus, even random GH levels < 1 µg/L do not rule out acromegaly. The two most used tests to assess biochemical activity in acromegaly are postglucose GH nadir (GHn) and IGF-1. GHn during an oral glucose tolerance test (OGTT) has been used as the standard method for acromegaly diagnosis (2), although we have recently reported suppression in low GH naïve-active acromegalic patients below the current consensus values (3,4). Furthermore, it has been reported that OGTT should ideally be evaluated in relation to age (5-8), gender (5,7) and body mass index (5,8). Actually, there has been several attempts to define GHn in normal patients using different assays, when cutoff values were lower than 0.3 (9,10). One report has suggested that random basal GH levels ≥ 5 µl in men, and ≥ 10 µl in women may possibly obviate the need of an OGTT suppression test, when IGF-1 is elevated (11). IGF-1 levels correlate with clinical features of acromegaly and are stable during the day (12). Studies have demonstrated that GHn and IGF-1 usually correlate well (13), although their relationship has not been tested in acromegalic populations with different GH outputs, and for different GHn thresholds.

Thus, we aimed at evaluating the correlation between GHn and IGF-1 levels in two different populations of clinically and biochemically naïve-active acromegalic patients: one with clearly elevated, and the other with low mean 24-h GH concentrations. We studied thirty-eight adults with untreated acromegaly (22 males, 16 females), as defined by the presence of clinical symptoms and exam results showing elevated age-adjusted IGF-1 values. Their ages ranged from eighteen to seventy-three years old. All patients had been enrolled in other reported protocols at the University of Michigan. All patients underwent frequent blood sampling for GH followed by an OGTT, when a sample for IGF-1 was also collected. A mean 24-h GH value of 4.3 µg/L was used as the threshold value to distinguish “high” and “low” GH study groups. This cutoff value was chosen because it represented the highest mean 24-h GH level in our studies of normal, healthy control adults (14). Fourteen patients were defined as “high” GH acromegalic patients (9 males, 5 females; mean age 40.6 ± 3.8 years) and twenty-four as...
"low" GH acromegalic patients (13 males, 11 females; mean age 46.3 ± 2.9 years). All patients had a MRI-identifiable pituitary adenoma. A GH-secreting pituitary adenoma was histologically and immunochemically identified in all cases (n = 38). None of the patients had renal or hepatic impairment, and none were treated with dopamine agonists, somatostatin analogues, GH receptor antagonists, estrogen or any medication that could potentially affect GH secretion.

Participants consumed a standard isocaloric hospital diet consisting of three meals and a bedtime snack during sample collection. Blood sampling for GH was performed every ten minutes for 24-h in all patients. Subsequently, an oral glucose load was given, followed by GH and glucose measurements every 10 min for two hours, between 07:00 and 09:00h (13 points in time), while patients were resting in a supine position. Serum GH was measured in duplicate using a chemiluminescent assay (Nichols Institute Diagnostics, San Juan Capistrano, CA, USA). Sensitivity of the GH assay reported by the manufacturer, and confirmed by our experience, was 0.01 µg/liter. Plasma IGF-1 was measured using a two-site immunoradiometric assay (Diagnostic System Laboratories, Webster, TX, USA) in all samples. Age-adjusted IGF-1 values were obtained from a previously published large multicenter study by Brabant e cols. (15), which employed the same assay used in this study. Z-score calculation required the average of age-specific standard deviation reference values, since reference data showed a slightly non-Gaussian distribution.

In order to address the relationship between GHn and IGF-1 levels, we subdivided the whole group of 38 patients according to both the group of origin ("high" or "low" GH group), and to GHn cutoff value of 1 µg/L. The relationship between logarithm-transformed GH levels and IGF-1 z-scores were analyzed by means of correlations using Spearman r in each analysis. As all patients from the "high" GH group had GHn higher than 1 µg/L, we had thus obtained two groups and two subgroups of patients according to these criteria, respectively: “high” GH group with GHn higher than 1 µg/L (N = 14), “low” GH group with any GHn value (N = 24), “low” GH group with GHn higher or equal than 1 µg/L (N = 11), and “low” GH group with GHn lower than 1 µg/L (N = 13).

The two studied groups did not differ significantly in terms of age, gender, and body mass index. The 24-h individual mean GH for the “high” and “low” GH patients ranged from 4.6 to 120.8, and 0.6 to 4.0 µg/L, respectively (P < 0.0001). There was a significant correlation between IGF-1 z-scores and mean 24-h GH for the entire group (r = 0.73, P < 0.0001). As shown in figure 1, significant correlations between GHn and IGF-1 z-scores were detected for both the entire group (r = 0.83, P < 0.0001) and the “low” GH group (r = 0.72, P < 0.0001).

![Figure 1](image.png)

**Figure 1.** Correlation between the logarithm-transformed postglucose growth hormone nadir (GHn) and insulin-like growth factor 1 (IGF-1) z-scores for the entire group (A; N = 38), and both “high” (B; N = 14), and “low” (C; N = 24) growth hormone groups.

However, GHn did not correlate with IGF-1 z-scores for the “high” GH group. As shown in figure 2, when the “low” GH group was subdivided into two
subgroups according to the cutoff value of 1 µg/L for GHn, those patients who showed suppression higher or equal to 1 µg/L showed significant correlation with IGF-1 z-score ($r = 0.80$, $P < 0.01$), while those who showed suppression lower than 1 µg/L did not show significant correlation with IGF-1 Z-score.

Altogether, these data show that the correlation between GHn and IGF-1 Z-scores, although significant for the entire group, is lost in the extremes of GHn, as demonstrated for the “high” GH group, and the “low” GH group with GHn lower than 1 µg/L. Interestingly, we have found GHn lower than 1 µg/L in more than 50% of our “low” GH acromegalic patients, and GHn lower than 0.4 µg/L in almost 30% of these patients. Moreover, we had two active female acromegalic patients, in which GHn is expected to be higher than in males, showing GHn levels as low as 0.03 µg/L and 0.07 µg/L. Accordingly, a mild residual disease activity in a subgroup of treated acromegalic patients with GHn < 1 µg/L and increased IGF-1 was recently shown (16).

In conclusion, besides OGTT being superfluous in the diagnosis of acromegaly in patients with active disease and “high” GH (3), and the difficulty in interpreting it in relation to age, gender and body mass index (4), there is an important lack of agreement between GHn and IGF-1 when GH output is too high or too low. In the first situation, repeated baseline GH levels, instead of GHn, may be of importance in follow-up alongside with IGF-1, due to possible saturation of GH receptors (12). More importantly, in the second situation, no matter how low GHn is, it is still not enough to claim biochemical inactivity, and suppressed GHn should not, therefore, be regarded as important whenever IGF-1 is high in acromegaly.

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