Bud dormancy in woody species is generally induced naturally in late summer and fall and is broken by low temperatures that occur during fall, winter, and spring. Temperature is the main factor that affects dormancy release. The efficiency of temperature to release buds from dormancy depends on its absolute value and on the range of temperatures in daily or longer cycles (Bennett, 1949; Couvillon and Erez, 1985b; Erez et al., 1979a, 1979b; Overcash and Campbell, 1955; Samish and Lavee, 1962). The “chill-unit model” developed by Richardson et al. (1975) takes into consideration the effects of various temperatures on the dormancy-breaking process; it usually has been useful in temperate climate zones (Shaltout and Unrath, 1983).

Bud dormancy status is estimated by measurement of growth rate or developmental stage of terminal, lateral, or isolated buds in excised shoots or small plants after buds have been exposed to environmental conditions favorable for growth (Couvillon and Erez, 1985a; Hatch and Walker, 1969; Landsberg, 1974; Latimer and Robitaille, 1981; Paiva and Robitaille, 1978a, 1978b; Shaltout and Unrath, 1983; Singh and Powell, 1978; Spiegel-Roy and Alston, 1979; Williams et al., 1979). To determine 50% flower bud growth, plants are subjected to temperatures of 16 to 24°C and 16 to 24 h of light. In that protocol, dormancy is generally considered over when 50% budbreak occurs within 14 to 21 days. The purpose of this study was to compare the different estimation indices for the length of the bud dormancy period in a wide range of apple cultivars and related Malus spp. growing under cold winter conditions. The CR of apple cultivars (Hauagge and Cummins, 1991a) and the seasonal variation in bud dormancy intensity (Hauagge and Cummins, 1991b) are considered in companion papers.

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Present address: IAPAR-Instituto Agronômico do Paraná, P.O. Box 2301, Curitiba, PR, Brazil 80001.

Abbreviations: BB, percent terminal budbreak; BDS, bud development stage; CR, chilling requirement; CU, chill unit; D50, number of days for 50% terminal budbreak; GDH, growing degree hours.
Arrangement of shoots was completely randomized, with each observation unit consisting of two shoots replicated three times in 1984–85 and four shoots replicated four times in 1985–86 and 1986–87. The data were subjected to regression analysis for the best fitted function, linear or quadratic. Only the last data point, the value of which was zero, was used in the regression for the bud development data. Regression was fitted for BB after removing zero and 100 values.

Results and Discussion

Field CU and GDH accumulation patterns were similar during the three experimental years (Figs. 1 and 2). No significant GDH were accumulated in Geneva during winter.

Intact shoots. The patterns for D50 and both BDS and BB after 21 days of forcing as a function of CU accumulation for ‘Anna’ and ‘Empire’ are represented in Fig. 3. A quadratic model fit better than a linear one for the D50 and BDS in all experimental years and all cultivars. Many genotypes had shown some tendency to shift the D50 to the right. This shift can be observed when good linear fit was obtained by plotting the inverse of D50 (Fig. 3F). However, not all cultivars showed this tendency (Fig. 3B), and usually very little or no improvement for the fitting was observed with these data. In this way, cultivar CU requirement values determined on BDS and D50 bases were interpolated from a 2nd-degree function. Either BDS or BB values were maintained at or near the 0 level, and then they rose abruptly in most cultivars (Figs. 3 G and H); however, in some cultivars with low depth of bud dormancy (Fig. 3D), this rise did not occur. In this situation, buds always developed within the forcing period.

Cultivar, linear, and quadratic components of regression, as well as all interactions, were significant for D50, BDS, and BB when regressed against CU accumulation (Table 1). Identical trends were also observed in the 1984–85 and 1986–87 seasons (data not shown). This result indicates that apple genotypes differ not only as to when changes in dormancy intensity take place, but also in the patterns and rates at which these changes proceed. This exemplifies the complexity of dormancy phenomena when one tries to consider a species as a whole. These particular genotype interactions imply that dormancy modelling must not be generalized or precision will be lost. These genotypic differences have implications for the adaptation of a particular genotype to diverse environmental conditions (Hauagge and Cummins, 1991b).

The means of the adjusted coefficient of determination for the quadratic fitting of D50 as a function of CU accumulation for all cultivars and *Malus* spp. were 0.79 ± 0.13 (0.56 to 0.96) and 0.82 ± 0.12 (0.64 to 0.99) for the 1985–86 and 1986–87 seasons, respectively. The values were similar to those obtained when the same data were fitted on the basis of the total GDH accumulation in the field and during forcing. Consequen...
quently, except for very high CR genotypes, GDH accumulated in the field can be ignored if the termination of dormancy is the only point studied. The probability for the F value for the model was ≤0.001 and, in most cases, ≤0.0001.

The means of the adjusted coefficient of determination for the quadratic fitting of BDS (1984-85 and 1985-86) and BB (1985-86) as a function of CU accumulated for all cultivars, and Malus spp. were 0.80 ± 0.13 (0.53 to 0.98), 0.67 ± 0.13 (0.41 to 0.94), and 0.66 ± 0.19 (0.42 to 0.93), respectively. In all cases, the model was significant at P ≥ 0.05.

The state of bud dormancy estimated by D50 is more precise than either BB or BDS because BB or BDS values varied abruptly from a minimum to a maximum range of CU accumulation (Fig. 3). D50 values varied in a much wider range of CU accumulation; consequently, more data points could be used in the regression. In addition, D50 values showed less variability and were less subjective. D50 values are especially interesting because they indicate the growth potential of buds when other indices show no changes. This characteristic may permit better understanding of development of dormancy throughout the season. Similar data were reported for peaches, where GDH was a more sensitive parameter for the estimation of bud dormancy state than was BB (Scalabrelli and Couvillon, 1986).

We observed that well-matured, dormant apple shoots (close to the time of natural leaf fall and afterward) can withstand up to 170 days under forcing conditions if proper care is taken—prevention of desiccation, weekly trimming of cutting bases, and constant running water at 12 to 15°C. Grafted plants of some apple cultivars under an alternating-temperature regime had also shown budbreak after being maintained 170 days under mild-temperature forcing conditions (Hauagge and Cummins, 1991a).

Table 1. Analysis of variance components for the linear and quadratic effects of the CU accumulation in three methods for the evaluation of bud dormancy status in apple cultivars in the 1985–86 season.

| Source       | df | Mean square | F value |
|--------------|----|-------------|---------|
| D50 Cultivar |    | 135.1       | 22.67***|
| D50 CU      |    |             |         |
| Linear      | 1  | 387.653     | 6202.96***|
| Quadratic   | 1  | 174.0       | 29.20***|
| Linear × cultivar | 97 | 51.7       | 8.67***|
| Quadratic × cultivar | 97 | 27.8       | 4.67***|
| Experimental error | 1939 | 5.9 |         |
| BDSa         |    |             |         |
| Cultivar     | 81 | 6.69        | 19.36***|
| CU           |    |             |         |
| Linear      | 1  | 475.74      | 1377.21***|
| Quadratic   | 1  | 270.84      | 784.05***|
| Linear × cultivar | 81 | 1.85       | 5.44***|
| Quadratic × cultivar | 81 | 0.90       | 2.60***|
| Experimental error | 1650 | 0.34 |         |
| BBb          |    |             |         |
| Cultivar     | 76 | 9899.6      | 19.66***|
| CU           |    |             |         |
| Linear      | 1  | 8465.968    | 1681.51***|
| Quadratic   | 1  | 2371.5      | 4.71***|
| Linear × cultivar | 76 | 2793.4     | 5.55***|
| Quadratic × cultivar | 76 | 1742.1     | 3.46***|
| Experimental error | 1346 | 503.5 |         |

a Estimated after 21 days of forcing.

**Significant at P = 0.05 or 0.001, respectively.

Nonetheless, a larger sample would improve precision at the time of maximum dormancy stage because a higher variability is observed at this point.

The total accumulated CU value needed for each taxon to reach an average specific level of D50 (20, 15, 10, or 5 days), BDS (0.25, 0.50, 1.50, and 2.00), and BB (25%, 50%, 75%, or 100%) was calculated by using fitted equations, and the values were correlated among themselves (Tables 2–4). D50 for

Table 2. Correlation coefficients between estimates for CU accumulation needed for D50 within 20, 15, 10, and 5 days of forcing in several apple genotypes in the 1985–86 and 1986-87 seasons.

| 1986–87  | 1985–86 Estimates† |
|---------|--------------------|
| Estimates | 20 days | 15 days | 10 days | 5 days |
| 20 days | 0.72*** | 0.72*** | 0.64*** | 0.37*** |
| n = 81  | n = 87  | n = 88  | n = 69 |
| 15 days | 0.70*** | 0.75*** | 0.72*** | 0.53*** |
| n = 81  | n = 87  | n = 88  | n = 69 |
| 10 days | 0.54*** | 0.58*** | 0.65*** | 0.54*** |
| n = 80  | n = 85  | n = 86  | n = 67 |
| 5 days  | 0.20**  | 0.27*   | 0.44*** | 0.52*** |
| n = 72  | n = 76  | n = 76  | n = 61 |

†n = Number of genotypes used in the comparison.

**Significant at P = 0.05, 0.01, or 0.001 or nonsignificant, respectively.

Table 3. Correlation coefficients between the number of accumulated CU required for terminal buds to reach 0.25, 0.50, 1.00, 1.50, and 2.00 BDS within 21 days of forcing in several apple genotypes in the 1984-85 and 1985-86 seasons.

| 1985–86  | 1984–85 Estimates† |
|---------|--------------------|
| Estimates | BDS 0.25 | BDS 0.50 | BDS 1.00 | BDS 1.50 |
| BDS 0.25 | 0.54  | 0.62  | 0.67  | 0.66  | 0.61  |
| n = 51  | n = 52  | n = 54  | n = 55  | n = 55 |
| BDS 0.50 | 0.58  | 0.66  | 0.70  | 0.69  | 0.63  |
| n = 55  | n = 58  | n = 61  | n = 62  | n = 63 |
| BDS 1.00 | 0.58  | 0.60  | 0.70  | 0.71  | 0.67  |
| n = 60  | n = 63  | n = 67  | n = 68  | n = 69 |
| BDS 1.50 | 0.56  | 0.60  | 0.69  | 0.73  | 0.70  |
| n = 63  | n = 66  | n = 69  | n = 70  | n = 72 |
| BDS 2.00 | 0.50  | 0.52  | 0.61  | 0.64  | 0.62  |
| n = 62  | n = 65  | n = 68  | n = 69  | n = 71 |

†All comparisons were significant at P = 0.001.

n= Number of genotypes used in the comparison.

Table 4. Correlations among the amount of accumulated CU needed for buds to reach specific levels of the parameters D50, BDS (after 21 days of forcing), and BB (after 21 days of forcing) used to estimate the state of bud dormancy of apple genotypes in the 1985–86 seasons.

| 1985–86  |
|---------|
| Criterion | D50 | BDS |
| 20 | 0.25 | 0.79* | 0.64 | 0.51 | 0.36 |
| 15 | 1.00 | 0.82 | 0.80 | 0.67 | 0.58 |
| 10 | 1.50 | 0.75 | 0.75 | 0.67 | 0.44 |
| 5  | 2.00 | 0.61 | 0.72 | 0.68 | 0.47 |

| BDS 0.25 | 0.89  | 0.86  | 0.71  | 0.42  | 0.89  |
| 0.87  | 0.90  | 0.83  | 0.78  |
| 0.87  | 0.90  | 0.83  | 0.78  |
| 0.87  | 0.90  | 0.83  | 0.78  |
| 0.87  | 0.90  | 0.83  | 0.78  |
| All correlations were significant at P = 0.001.

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15 days and BDS for 1.50 gave the highest correlation for independently obtained D50 (Table 2) and BDS (Table 3) value estimates. The correlation among independent estimations of either BDS and D50 showed some tendency to decrease in both sides (Tables 2 and 3) around the maximum values. BDS = 1.5 and D50 = 15 values also correlated well when comparisons were made within the same year (Table 4), as well as with BB = 50%. The BDS mean value for all clones was determined to be equal to 1.66 ± 0.44 when calculated for CU values obtained from D50 equations at the 15th day of forcing.

According to the concept of Shaltout and Unrath (1983), dormancy is broken when further budbreak is not induced by additional CU. This may be precise, but it is difficult to determine experimentally. For practical purposes, however, dormancy should be considered to have been broken when no more visible symptoms of “delayed foliation” can be observed. This stage seems to be when budbreak occurred within 21 and 23 days (20C GDH equivalent) from the initial forcing day in ‘Rome Beauty’ and ‘Golden Delicious’, respectively (modified from Tabuenca and Jiménez, 1984). The values of these characteristics are close to those generally adopted to describe the end of the bud dormancy period.

Summarizing, the indices BB = 50%, BDS = 1.5, and D50 = 15 correlated well among themselves, and they are close to values reported in the literature for the end of apple bud dormancy period.

Shoots from which terminal buds were removed. Decapitated shoots have been used to estimate the length of the bud dormancy period in Pyrus spp. (Spiegel-Roy and Alston, 1979) and Malus spp. (Petropoulou, 1985). The uppermost bud of decapitated shoots will always break within a relatively short period, independently of sampling date (Paiva and Robitaille, 1978a; Williams et al., 1979). The dormancy intensity pattern of the uppermost bud may have a trend similar to that of terminal buds in intact shoots, but much less accentuated (Hauagge and Cummins, 1991 b). The direct use of these data to create an index for-dormancy development was difficult because the differences in dormancy intensity in many genotypes are not clear. Considerable difference can be observed between Petropoulou’s 1985 estimation of dormancy termination using decapitated shoots and that estimated by us (Fig. 4).

CU accumulation until the time when the ratio between the D50 in decapitated shoots and the D50 in intact shoots (D50R) is equal to one is relatively close to the other estimations of CR to terminate bud dormancy reported here (BB = 50%, BDS = 1.5, D50 = 15). These values were determined by fitting D50R obtained on five sampling dates into quadratic or linear equations (Hauagge, 1988). However, D50R indices of the end of dormancy (accumulated CU to D50R = 1) are not as closely related to BB, BDS, and D50 indices as they are related to each other. For instance, correlations between D50R value estimates and BDS = 1.5 (1984–85), D50 = 15 (1985–86), and D50 = 15 (1986–87) were 0.46, 0.48, and 0.56, respectively. All val-

Table 5. Correlation coefficients between bud CR to break dormancy in apple genotypes and the dormancy intensity of the same genotypes at different sampling dates during summer and winter. CR was considered to be equal to CU accumulated when D50 was equal to 15 days. Bud dormancy intensity was estimated by D50 in intact shoots, decapitated shoots, and by the ratio between decapitated and intact shoots at the time of sampling (1986-87 season).

| Sampling date or accumulated CU | Intact shoots | Decapitated shoots | Decapitated/ intact shoots |
|--------------------------------|---------------|--------------------|--------------------------|
| 14 July                         | 0.18NS (n=79) | 0.33** (n=85)      | -0.25NS (n=24)           |
| 1 Sept.                         | 0.11NS (n=24) | 0.21* (n=80)       | 0.24NS (n=44)            |
| 289                             | 0.32** (n=61) | 0.38** (n=84)      | -0.36** (n=49)           |
| 741                             | 0.51** (n=94) |                     |                         |
| 880                             | 0.69** (n=94) | 0.66** (n=88)      | 0.51** (n=88)            |
| 1098                            | 0.72** (n=94) |                     |                         |
| 1193                            | 0.67** (n=94) | 0.68** (n=85)      | 0.19* (n=85)             |
| 1348                            | 0.36** (n=48) |                     |                         |
| 1366                            | 0.76** (n=45) |                     |                         |
| 1592                            | 0.77** (n=20) |                     |                         |

*NS: Nonsignificant or significant at P = 0.05 or 0.01, respectively.

Fig. 4. Relationship between CR index using decapitated shoots estimated by Petropoulou (1985) in Kent, England, and CU required to terminate terminal bud dormancy in Geneva, N.Y.

Fig. 5. Relationship between genotype CU accumulation required to achieve 50% BB at the 15th day of forcing and D50 on two sampling dates of the same genotypes.

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ues were significant at $P = 0.001$ and involved 62, 72, and 82 comparisons, respectively. The physiological significance of differences between the estimates is unknown. They reflect genotypic differences in relation to the development of the uppermost bud in response to decapitation. It is possible that a more detailed study on development of budbreak on intact shoots and decapitated ones might increase understanding of dormancy development and termination. In apple, the most obvious symptom of incomplete dormancy release under subtropical conditions is the absence (or very long delay) of lateral vegetative budbreak.

Correlations between CU accumulation required to D50 = 15 days and D50 values obtained at several sampling dates for the same genotypes are shown in Table 5. In general terms, terminal bud dormancy intensity during summer is not correlated with the length of bud dormancy observed either in intact shoots (within the cultivars that showed budbreak during the forcing period) or decapitated ones. The relationship improves afterward, stabilizing at ≈880 CU. This relationship may be helpful for a preliminary screening for CR in a large population. Such screening would be almost impossible by traditional methods, in which frequent sampling throughout the winter is required. For instance, a general idea about the length of bud dormancy (CR) could be derived by sampling at 880 CU accumulation (Fig. 5), but this relationship is almost nonexistent on the 14 July 1984 sampling.

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