Comparative Distribution and Metabolism of Xylem-Borne Amino Compounds and Sucrose in Shoots of *Populus deltoides*

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ABSTRACT

The transport and metabolism of xylem-borne amino compounds and sucrose were investigated in rapidly growing shoots of cottonwood (*Populus deltoides* Bartr. ex Marsh.). ¹⁴C-labeled glutamine, threonine, alanine, glutamic acid, aspartic acid, and sucrose were applied to the base of severed stems for transport in xylem. Distribution and metabolism of the compounds were followed with autoradiography, microautoradiography, and radioassay. Three utilization patterns were observed: (a) little alanine and sucrose was transported to the laminae of either mature leaves or developing leaves. These compounds were taken up from xylem free-space and utilized in adjacent tissue; (b) threonine also did not move into mature leaves but was translocated to developing leaves or utilized in the stem; (c) glutamic acid and aspartic acid were transported directly into the laminae of mature leaves via the xylem. Relatively less ¹⁴C was retained in stems compared to the other compounds.

Metabolism of the test compounds also differed considerably. ¹⁴C from amino acids moved primarily into organic acids and protein. The ¹⁴C from sucrose was widely distributed among the chemical fractions, with a high percentage found in structural carbohydrates. Clearly, cottonwood stems contain efficient uptake and transfer systems that differentiate among various compounds moving from root to shoot in xylem.

The transport of nitrogenous compounds from root to shoot in xylem and the transfer of these compounds from xylem to phloem in the shoot has been investigated with intensive studies on legumes using modern techniques for the efficient collection of xylem and phloem sap (20). Such studies have examined in detail the movement of nitrogen from root to shoot and the cycling of both nitrogen and carbon within the plant (12, 16, 18, 29). Similar intensive studies have not been conducted on nonleguminous plants or trees. Most studies on the transport of nitrogen in trees have examined either naturally occurring products in xylem sap (2, 6, 24), followed changes in nitrogen transport after different fertilizer treatments (10, 28), or studied stem extracts of nitrogen storage products (7, 15, 23).

The two amides, glutamine and asparagine, are major transport compounds in trees and many other woody plants (24). These amides, as well as some other amino acids, move readily in both xylem and phloem to developing tissues (16). However, initial distribution within the plant differs considerably among plant species. For example, from 30 to 40% of the glutamine fed to severed lupin stems passed through the stem xylem and into mature leaves (12). In contrast, glutamine fed to cottonwood was largely removed from the xylem in the stem (5). Several other amino acids commonly found in xylem sap also differ widely in transport characteristics. For example, aspartic acid fed to the

xylem of severed lupin stems accumulated primarily in the leaf mesophyll while arginine accumulated in stem and leaf vascular tissue (12). Metabolism of xylem-borne amino acids can also differ both during and after transport. Sharkey and Pate (26) found that asparagine was largely transferred unchanged from xylem to phloem while aspartic acid was metabolized to several other products before loading into the phloem. Our objective was to follow the transport, uptake, and metabolism of sucrose and the major amino acids found in cottonwood xylem sap after application of these ¹⁴C-labeled compounds to the bases of severed stems.

MATERIALS AND METHODS

Plant Material. Cottonwood plants (*Populus deltoides* Bartr. ex Marsh.) were grown from seed in a controlled environment room under conditions previously described (5). However, in the present series of experiments, the plants had eight mature leaves (LPIs¹ 7–14) attached during treatment compared to three or seven mature leaves per plant in previous experiments. (5).

Application of Radioactive Compounds. Early in the treatment day, plants were removed from the growth room and preconditioned as before (5). After 1 h, the leaf at LPI 15 was removed and the plants were severed under water at the base of the internode subtending LPI 15. The severed plants were transferred to 50-ml beakers containing artificial xylem sap (5 mol m⁻³ KCl, 0.4 mol m^{-3} malic acid, adjusted to pH 5.4 with KOH) and placed in the treatment chamber. The plants were next transferred to vials containing 0.9 MBq of either [U-14C]glutamine, -glutamic acid, -aspartic acid, -alanine, -threonine, or -sucrose per ml of artificial xylem sap. Each plant was fed 1 ml of radioactive solution (approximately 15 min) and then transferred back to unlabeled solution for the remainder of the translocation period. A single plant was treated for either 0.5, 1, or 3 h (feeding time plus chase with unlabeled solution) to provide a translocation-metabolic time series for each compound. The time series were not replicated because previous work (5) showed that uptake and distribution patterns were consistent from plant to plant, the extraction procedures were reliable (3), and distribution of ¹⁴C in different tissues and chemical fractions across the time series provided a check on large errors.

Plant Harvest, Autoradiography, and ¹⁴C Analyses. At the end of the translocation period, internode, petiole, and leaf segments were excised, frozen in liquid N₂, freeze-dried, pressure infiltrated with Spurr's low viscosity resin, sectioned, and prepared for microautoradiography (32). The remaining leaves and stem segments were pressed, quick-frozen in dry ice-liquid N₂ vapor, freeze-dried, and autoradiographed with Kodak Industrex X-ray

¹ Abbreviations: LPI, leaf plastochron index; GAB, gamma-aminobutyric acid.

After freeze-drying and autoradiography, samples of lamina and stem (5-15 mg) were extracted with methanol:chloroform:water (12:5:3, v:v:v) and separated into seven chemical fractions with a combination of solvent, ion-exchange, and enzymic techniques (3). The seven fractions were CHCl₃ (lipids and pigments soluble in chloroform), protein (pronase extract of the pellet), starch (Clarase and Diazyme, Miles Laboratories², extract of the pellet), residue (tissue remaining after all extracts), sugar (neutral fraction from the ion exchange columns), organic acids (6 M HCOOH eluate from the anion column), and amino acids (4 м NH4OH eluate from the cation column). Total recovery of ¹⁴C (sum of the seven fractions) averaged about 95% based on NCS analysis of the tissue. The individual amino acids and organic acids were separated with TLC, scraped from the thinlayer plates into scintillation vials, and solubilized in NCS for ¹⁴C determinations.

RESULTS

Localization of ¹⁴C in Whole Leaves and Stem Internodes. The movement of radioactive amino acids into leaves after xylemfeeding cottonwood plants resulted in three general distribution

² Mention of trade names is for the convenience of the reader and does not constitute endorsement by the United States Department of Agriculture Forest Service.

patterns. (a) Alanine did not accumulate in mature leaves or in young leaves (Fig. 1A; Table I). (b) Threonine also did not accumulate in mature leaves but rapidly accumulated throughout the lamina in young developing leaves (Fig. 1B). (c) ¹⁴C from glutamic and aspartic acid rapidly accumulated in the mesophyll of mature leaves (Fig. 1C).

The accumulation patterns of aspartic acid or glutamic acid indicated both xylem and phloem transport. In mature leaves, some loading was apparent in the major laterals and midvein after 1 h (Fig. 1C); however, most of the ¹⁴C remained in the interveinal mesophyll. In developing leaves, initial accumulation of glutamic or aspartic acid (after 0.5 h) was primarily in lamina tips and marginal glands indicating xylem transport. Labeling of leaf glands was most noticeable in aspartic acid-treated plants (LPIs 0–2, Fig. 1C). With longer treatment time (3 h), ¹⁴C concentration increased in the smaller veins and mesophyll.

The distribution of ¹⁴C in the leaves after xylem-feeding sucrose differed from that obtained with amino acids. The major veins of all mature leaves were rapidly labeled (Fig. 1D); the ¹⁴C from sucrose was confined to the veins and did not move into the interveinal mesophyll even after 3 h. Initially (after 0.5 h), young developing leaves contained little ¹⁴C, and that present was confined to the vascular tissue. At 3 h, vein labeling had increased and some ¹⁴C had accumulated in the interveinal mesophyll.

Specific activity of ¹⁴C in the laminae quantitatively confirmed the autoradiographic patterns (compare Fig. 1 with Table I). For example, the concentration of ¹⁴C in leaves at LPI 8 reflected the



FIG. 1. Autoradiographs of whole cottonwood leaves after xylem-feeding different amino acids and sucrose. The plants were treated for 15 min (0.9 MBq ¹⁴C), harvested 1 h after the ¹⁴C-labeled compounds were applied, freeze-dried, pressed, and autoradiographed. A, Alanine-; B, threonine-; C, aspartic acid-; and D, sucrose-treated plants. Each series shows a recently matured leaf and five or six developing leaves plus the terminal bud and elongating internodes. See threonine (B) for the LPI numbering sequence. The lamina of LPI O was 2 cm long.

Table I. Specific Activity of ¹⁴C in Laminae and Stems of Cottonwood after Xylem-Feeding Radioactive Amino Acids and Sucrose

Plants were fed 0.9 MBq ¹⁴C-amino acids or sucrose through the xylem of the severed stem and then transferred to a nonradioactive solution for the indicated translocation times. Specific activity (Bq/mg dry wt plant material) is the sum of ¹⁴C recovered in each of seven chemical fractions separated during chemical analysis of the different plant tissue. Stem segments LPI 4, 9, and 12 consisted of the node and internode directly subtending the leaf of that age and contained both xylem and phloem.

	¹⁴ C Specific Activity at Following Times (h)														
	Laminae					Stems									
	LPI 4				LPI 8			LPI	4		LPI 9			LPI 12	
	0.5	1	3	0.5	1	3	0.5	1	3	0.5	1	3	0.5	1	3
	Bq/mg dry wt														
Glutamic acid	274	412	504	75	228	108	272	900	1504	813	2005	516	921	1539	1275
Aspartic acid	200	602	573	543	306	328	262	295	319	462	742	287	346	922	438
Alanine	71	88	256	13	33	52	884	992	589	797	1799	1036	1768	3711	2113
Threonine	103	481	480	6	14	21	723	801	760	2090	1622	1295	2862	1757	2364
Sucrose	66	68	255	94	163	79	311	454	1666	1226	1108	2342	1118	1161	1861

Table II. Specific Activity of ¹⁴C in Developing Leaves of Cottonwood after Xylem-Feeding ¹⁴C-Labeled Aspartate and Threonine

Specific activity is expressed as Bq/mg dry wt of lamina tissue. Leaves were freeze-dried, ground, subsampled, and solubilized in NCS to determine the ¹⁴C concentration. Lamina dry weights are averages obtained from 16 plants grown under the same experimental conditions. Dry weights times specific activity will give the total ¹⁴C present in the different-aged leaves.

	•	¹⁴ C Specific Activity at Following Times (h)								
LPI	Lamina Dry Wt	Asj	partic a	ncid	Threonine					
		0.5	1	3	0.5	1	3			
	mg	Bq/mg dry wt								
0	9.4	9	195	454	122	507	890			
1	16.6	45	348	487	135	436	868			
2	31.5	65	460	506	216	570	848			
3	54.5	197	802	696	189	649	786			
4	86.5	315	854	795	131	413	456			
5	125.5	389	988	850	40	211	212			
7	164.8	506	632	607	7	15	14			
9	196.7	664	443	574	4	15	6			
12	174.1	186	151	196	2	5	17			

more rapid movement of glutamic and aspartic acid into mature laminae compared to alanine and threonine. Activity in the leaf at LPI 4 was fairly high after treatment with all of the compounds except alanine and sucrose and generally increased with time. Specific activities in different-aged stem segments show that more alanine, threonine, and sucrose accumulated throughout the lower stem—LPIs 9 and 12—compared to glutamic and aspartic acid (Table I).

Comparing specific activities of aspartic acid and threonine after 0.5 h in a series of leaves differing widely in age showed that the concentration of aspartic acid was directly related to leaf size and/or transpiration rate (Table II). ¹⁴C increased in the youngest leaves of aspartic acid-treated plants only after prior movement into mature leaves and retransport in the phloem. In contrast, little threonine moved into the older leaves but rapidly accumulated in young developing leaves.

Microautoradiography of Stem and Leaf Tissues. Microautoradiography showed that alanine and threonine uptake by the stems was similar. ¹⁴C rapidly accumulated in the xylem parenchyma, rays, and differentiating cells of the cambial zone in the lower stem (Fig. 2, A and B). Concentrations were particularly high in the metaxylem parenchyma of traces serving mature leaves on the lower stem, in ray cells, and in vessel-associated parenchyma cells (Fig. 2C). Little ¹⁴C accumulated in phloem of the lower stem. However, in the upper stem (internode LPI 7), considerable [¹⁴C]threonine had accumulated in the phloem primarily in traces serving recently matured and expanding leaves (Fig. 2D). At LPI 3, ¹⁴C from threonine was concentrated in phloem of both the internode and petiole (Fig. 2E), although some scattered activity occurred throughout these tissues (Fig. 2F).

2F). ¹⁴C from aspartic and glutamic acid was widely scattered throughout the stem tissue (Fig. 3A), with accumulations in outer pith, metaxylem parenchyma, rays, cambium, and phloem. In the upper stem (internode at LPI 3), most ¹⁴C was in the phloem (Fig. 3B). Transfer to phloem occurred in recently matured leaves. Microautoradiography of a major lateral vein and petiole of a leaf at LPI 8 after 1 h showed little ¹⁴C from aspartic acid present in the xylem but high concentrations in the phloem (Fig. 3, C-E). The highest ¹⁴C concentrations were in phloem of the dorsal bundles of the petiole, indicating that loading and retransport was taking place primarily in the apical portion of the lamina, an area of relatively older tissue.

Sucrose accumulated in differentiating cells of the secondary xylem of the lower stem (Fig. 4A), and in differentiating sclerenchyma cells located in the protoxylem region of the oldest leaf traces (Fig. 4A, arrow). In contrast to the amino acids, little sucrose was found in metaxylem parenchyma, older ray tissue, cambium, or phloem. Farther up the stem, in primary tissue of the internode at LPI 5, most ¹⁴C was found in traces serving leaves at LPIs 5 and 4, leaves with substantial transpiration (Fig. 4B). Within these traces, ¹⁴C was concentrated in the walls of differentiating metaxylem vessels and in vessel-associated parenchyma (Fig. 4B). Some ¹⁴C was present in the phloem, presumably the result of retransport from mature leaves lower on the stem.

In the petiole of a mature leaf (Fig. 4C), ¹⁴C accumulated in differentiating cells of the xylem, phloem, and phloem fibers. Much of the ¹⁴C apparently was incorporated into developing cell walls (Fig. 4D). In addition, areas of accumulation in the mature phloem indicate retransport of imported sucrose. Microautoradiographs of a petiole at LPI 6 (Fig. 4, E and F) showed accumulation of ¹⁴C in xylem parenchyma and in vessel cell walls. Little accumulated in phloem fibers because secondary wall development had not yet begun in the LPI 6 leaf.



FIG. 2. Microautoradiographs of cottonwood internode and leaf transections after xylem-feeding labeled alanine or threonine. A–C, treatment: $[U^{-14}C]$ alanine, 0.5 h (15-min pulse, 15-min chase); exposure, 8 d. A, Internode at LPI 12 showing short-term accumulation in metaxylem parenchyma (Mx), rays, developing secondary xylem, and cambium (× 65). B, Section from the upper left quadrant of A showing accumulation of ¹⁴C in cambium (C), young developing rays (upper arrow), mature rays (lower arrow), and developing secondary xylem (Sx). Little ¹⁴C accumulated in the phloem (P) (× 163). C, Section from lower right quadrant of A showing heavy accumulation of ¹⁴C in metaxylem parenchyma, in ray parenchyma (arrow), and in vessel-associated parenchyma (dart) (× 163). D–F, treatment: $[U^{14}C]$ threonine, 1 h (15-min pulse, 45-min chase); exposure, 8 d (D) and 40 d (E). D, Internode at LPI 7 showing high levels of ¹⁴C in cambium, rays, and metaxylem parenchyma in old leaf trace (left) and phloem loading in an adjacent younger trace (right, arrow) (× 130). E, Mid-petiole of a leaf at LPI 3 showing ¹⁴C concentrated in phloem and cell walls of developing xylem (× 58). F, Enlargement of boxed area of G. Note distribution of [¹⁴C]threonine throughout the petiolar tissue (× 163).

Distribution of ¹⁴C in Different Chemical Fractions of Leaves and Stems. In laminae at both LPIs 4 and 8, ¹⁴C from the labeled amino acids moved rapidly into the organic acid fraction. For example, up to 40% of the ¹⁴C was found in the organic acids within 0.5 to 1 h when aspartic acid was applied (Table III). The amino acids were also rapidly incorporated into protein, particularly in the developing leaf (LPI 4). ¹⁴C incorporated into protein in LPI 4 ranged from 27% (aspartic acid) to 70% (threonine). Alanine differed from the other amino acids in the small amount of ¹⁴C converted to organic acids and the large amount recovered



FIG. 3. Microautoradiographs of cottonwood internode and leaf transections after xylem-feeding labeled glutamic acid and aspartic acid. A and B, treatment: $[U^{-14}C]$ glutamic acid, 1 h (15-min pulse, 45-min chase); exposure, 10 d. A, Internode at LPI 8 showing ¹⁴C scattered throughout the stem with some accumulation in outer pith (Pi), metaxylem parenchyma (Mx), rays, and cambium (C) with less in the phloem (P) (× 130). B, Internode at LPI 3 showing ¹⁴C located mainly in the phloem (P) (× 163). C and D, Treatment: $[U^{-14}C]$ aspartic acid, 1 h (15-min pulse, 45-min chase); exposure, 40 d. C, a Major lateral vein of a mature leaf at LPI 8 showing ¹⁴C concentrated in the adaxial phloem (arrow); other phloem bundles were unlabeled (× 205). D, Mid-petiole of a mature leaf at LPI 8 showing accumulation of ¹⁴C in phloem. Although ¹⁴C was present in phloem of all vascular traces, the highest concentrations occurred in the dorsal bundles (boxed area and below) (× 52). E, Enlargement of the boxed area of D showing ¹⁴C concentrated in the phloem but little in the xylem and other tissues (× 130).



FIG. 4. Microautoradiographs of cottonwood internode and leaf transections after xylem-feeding labeled sucrose. A–F, treatment: $[U-I^4C]$ sucrose, 1 h (15-min pulse, 45-min chase); exposure, 5 d (C and E), 10 d (A), and 20 d (B). A, Internode at LPI 7 showing ¹⁴C from sucrose concentrated in the differentiating xylem (darts). At high magnification, much of the ¹⁴C appeared to be associated with or incorporated in the xylem cell walls. Differentiating sclerenchyma cells in the region of obliterated protoxylem of older leaf traces also accumulated ¹⁴C (arrow) (× 65). B, Internode at LPI 5 showing ¹⁴C concentrated in metaxylem (arrows) of traces serving leaves at LPIs 5 and 4; *e.g.* right trace of LPI 5 (5R) and left (4L) and central (4C) traces of LPI 4 (× 65). C, Mid-petiole of a leaf at LPI 9 showing ¹⁴C concentrated in differentiating xylem and phloem fibers (× 52). D, Enlargement of boxed area of C. Xylem (X), phloem fibers (F), and vessel-associated parenchyma cells (arrow). Note weak ¹⁴C loading of phloem (P) (× 150). E, Mid-petiole of a leaf at LPI 6 with ¹⁴C located mostly in the xylem (× 52) F, Enlargement of boxed area of E. Differentiating metaxylem vessels (arrow) and vessel-associated parenchyma are most heavily labeled (× 150).

Table III. Distribution of ¹⁴C among Major Chemical Fractions in Cottonwood Laminae after Xylem-Feeding Different Radioactive Amino Acids and Sucrose

¹⁴C-labeled amino acids and sucrose were fed through the xylem of the lower stem for 15 min and then chased with unlabeled solution. Plants were harvested at the times shown, freeze-dried, and subsampled (5-15 mg). The subsample was then separated into seven chemical fractions. One plant was used for each treatment time and each amino acid. LPI 4 was a developing leaf about one-half expanded. LPI 8 was a fully expanded and exporting leaf. Percentages are based on the total ¹⁴C recovered in the seven chemical fractions (see sucrose for these fractions). The petiole and midvein were removed before lamina analyses. CHCl₃, chloroform-soluble fraction, primarily pigments and lipids; OA, organic acids; AA, amino acids; Other, sum of all fractions not actually listed that contained minor amounts of ¹⁴C.

	Per Cent Total ¹⁴ C Recovered at Following Times (h)								
Chemical Fractions		LPI 4		LPI 8					
	0.5	1	3	0.5	1	3			
D			Gluta	amine		• •			
Protein	18	19	41	6	6	20			
ŬA.	32	28	27	33	32	35			
AA	40	42	18	44	48	33			
Other	10	11	14	18	14	12			
		(Glutan	nic acid	l				
Protein	11	16	29	6	8	20			
OA	20	22	16	16	22	18			
AA	62	55	40	74	65	52			
Other	7	7	15	4	5	10			
	Aspartic acid								
Protein	23	29	27	8	14	19			
OA	31	36	41	33	42	51			
AA	42	31	20	56	41	25			
Other	4	4	12	3	3	5			
			Alo	nina					
CHCh	18	0	16	6	12	12			
Protein	38	36	41	12	11	15			
Sugar	50 A	30	5	10	19	20			
O A	13	18	18	10	17	20			
A A	24	29	15	50	30	30			
Other	3	4	5	3	3	3			
Protoin	25	40	1 nre	onine	25	AE			
	33	00	/0	12	33	43			
	50	$\frac{2}{2}$	12	39	12	0			
Other	5	8	15	20	22	32			
	•		Suc	rose					
CHCl ₃	9	6	13	4	3	6			
Protein	25	19	27	13	10	14			
Starch	6	5	5	2	1	4			
Kesidue	0	0	0	3	2	1			
Sugar	54	40	29	28	57	49			
	11	01	14	12	21 4	1/			
<u>AA</u>	<u>у</u>	ō	0	ō	O	3			

in the CHCl₃ fraction. ¹⁴C from sucrose was also rapidly incorporated into protein and was widely distributed throughout the other chemical fractions in developing laminae.

The distribution of ¹⁴C among the chemical fractions in stem

Table IV. Distribution of ¹⁴C among Major Chemical Fractions in Cottonwood Stems after Xylem-Feeding Different Radioactive Amino Acids and Sucrose

Plants treated as in Table III. The stem segments given below comprised the node and internode subtending leaves at LPIs 4, 9, and 12 and contained both xylem and phloem.

	Per Cent Total ¹⁴ C Recovered at Following Times (h)										
Chemical Fractions		LPI 4	ļ	LPI 9			LPI 12				
	0.5	1	3	0.5	1	3	0.5	1	3		
					%						
				Gl	utam	ine					
Protein	6	8	24	2	4	14	2	3	8		
OA	22	34	35	25	28	27	22	22	26		
AA	68	54	34	68	64	52	72	72	53		
Other	4	4	7	5	4	7	4	3	13		
	Glutamic acid										
Protein	6	7	11	5	7	21	10	6	16		
OA	22	15	17	14	13	13	15	12	10		
AA	68	74	64	77	74	55	67	76	65		
Other	4	4	8	4	6	11	8	6	9		
				Asp	artic	acid					
Protein	11	11	14	7	14	19	7	16	19		
OA	30	40	33	28	10	24	20	18	18		
AA	52	44	47	61	70	48	68	57	50		
Other	7	5	6	4	6	9	5	9	13		
				А	lanin	e					
CHCl ₃	8	4	13	4	3	7	3	3	5		
Protein	25	35	40	24	31	33	18	29	27		
OA	15	27	14	8	19	16	5	10	10		
AA	44	20	18	54	32	27	66	44	43		
Other	8	14	15	10	15	17	8	14	15		
				Th	reoni	ne					
Protein	17	48	65	13	40	66	7	29	53		
AA	78	46	25	84	55	26	91	67	42		
Other	5	6	10	3	5	8	2	4	5		
				S	ucros	e					
CHCl ₃	6	6	4	4	5	6	6	5	4		
Protein	23	24	20	16	17	19	16	18	22		
Starch	12	14	13	8	9	16	9	10	8		
Residue	12	16	31	18	19	46	17	23	24		
Sugar	21	19	13	35	36	6	33	28	14		
OA	14	14	14	7	6	2	8	6	6		
AA	12	7	5	12	8	5	11	10	22		

segments differed from that found in laminae (Table IV). For example, relatively large amounts of ¹⁴C remained in amino acids for most of the experimental period. Threonine and alanine, two amino acids heavily accumulated in the stem, were rapidly incorporated into protein. ¹⁴C again accumulated in organic acids, particularly when glutamine and aspartic acid were applied. Sucrose was removed from the transpiration stream and utilized in a variety of storage and metabolic functions, as shown by the wide distribution of ¹⁴C among the different chemical fractions. Particularly important was the utilization in structural development (residue fraction) in the older stem segments.

Metabolic Changes within the Amino Acid and Organic Acid Fractions. Xylem-fed glutamine was rapidly deaminated in both leaves and stems and the resulting glutamic acid partially converted to GAB (Table V). Glutamic and aspartic acids were

Table V. Distribution of ¹⁴C among Individual Amino Acids from Cottonwood Laminae and Stems after Xylem-Feeding Different Radioactive Amino Acids and Sucrose

The amino acids were separated from other water-alcohol-soluble compounds with ion exchange and then isolated with TLC. Amino acid standards were spotted with the experimental solution on the TLC plates. After separation and visualization, the standard spots with the unknown ¹⁴C were scraped into scintillation vials for counting. Data shown are from the 1-h treatment. Percentage changes over time (0.5-3 h) show that ¹⁴C decreased in the fed amino acid and increased in other amino acids. Threonine was not determined in Leaf LPI 8 because there was not enough ¹⁴C in the AA fraction for reliable counts after separation. All amino acids analyzed are listed under sucrose.

	Per Cent Total Amino Acids									
Amino Acids	Lam	inae		Stems						
	LPI 4	LPI 8	LPI 4	LPI 9	LPI 12					
			%							
			Glutamir	ne						
Glu	45	44	58	50	41					
Asp	2	3	7	8	3					
Gln	1	2	6	20	19					
GAB	50	47	20	17	32					
Other	2	4	9	5	5					
	Glutamic acid									
Glu	53	47	69	42	34					
Asp	2	2	9	7	6					
Gln	1	1	5	11	19					
GAB	42	49	13	37	33					
Other	2	1	4	3	8					
	_	/	Aspartic a	cid						
Glu	2	2	9	13	16					
Asp	96	93	78	68	58					
GAB	0	1	8	13	18					
Other	2	4	5	6	8					
			Alanine	•						
Glu	46	19	44	38	24					
Asp	3	0	27	23	10					
Ala	24	67	9	10	25					
GAB	21	8	11	21	32					
Other	6	6	9	8	9					
			Threonir	ne						
Thr	80	_	94	94	95					
Other	20	_	6	6	5					
			Sucross							
Chu	21	18	34	37	18					
Ciù Ace	21	10	۲ د ۲	12	10					
Asp	, 1	17	1	12	1					
Asii	15	10	2	2	3					
Sar	15	20	22	20	16					
SCI Th-	22	5	23	20 2	2					
	0 17	12	10	17	35					
GAB	6	7	21	14	20					
U/U	v	,	~ .							

slowly metabolized to other amino acids in both leaves and stems. Glutamic acid was primarily converted to GAB in leaves, and to aspartic acid and glutamine in stems. Up to 30% of the tracer fed as [¹⁴C]glutamic acid was recovered as [¹⁴C]glutamine after 3 h in the stem at LPI 12. Glutamic acid and glutamine are readily interconverted in cottonwood stems although glutamic acid production is favored. Little aspartic acid was converted to other amino acids in leaves. More than 90% of the ¹⁴C was still in aspartic acid after 3 h. In stems, however, aspartic acid was converted to glutamic acid and GAB. After 3 h (data not shown), only 30% of the ¹⁴C was still in aspartic acid, while glutamic acid and GAB contained 20% and 36%, respectively. Alanine was rapidly converted to glutamic acid and aspartic acid in both young leaf and stem tissues. In contrast to other amino acids, little threonine was converted to either organic acids or to the other amino acids analyzed (Tables III–V).

The small amount of ¹⁴C from sucrose that was incorporated into the amino acid fraction of young leaves and stems (Tables III and IV) was found in several amino acids (Table V). The high percentage of ¹⁴C in serine is interesting because none of the amino acids contributed significant ¹⁴C to serine in any tissue or at any treatment time.

The percentage distribution of ¹⁴C among the organic acids often differed between leaves and stems after treatment with a particular amino acid (Table VI). For example, malic acid was lightly labeled in laminae but heavily labeled in stems after glutamine treatment. In both leaves and stems, malic acid and succinic acid contained high levels of ¹⁴C after aspartic acid, alanine, and glutamic acid treatment. Citric acid (from alanine, aspartic acid, and sucrose) and 2-ketoglutaric acid (from glutamic acid and alanine) also contained considerable ¹⁴C. Phosphorylated compounds (origin of the TLC plate) were heavily labeled after certain treatments (glutamine and alanine, LPI 8; alanine and aspartic acid, stems; sucrose, both leaves and stems).

DISCUSSION

The nitrogen requirements of different plant parts vary widely, depending on the developmental stage of the particular organ and of the whole plant. Young developing leaves or flowers have low transpiration rates and therefore must receive most of their nitrogen through the phloem (16, 29). Nitrogen in the phloem of developing leaves originates in mature leaves and from xylem to phloem transfer in the stem (4, 18).

Uptake from the Xylem and Transfer to the Phloem. Initial uptake from the xylem free-space takes place in the metaxylem, vessel-associated parenchyma, and ray parenchyma. These tissues form the major stem storage pools for both long-term storage in the fall (14, 25) and short-term storage involved in primary xylem to secondary xylem, xylen-to-phloem, and phloem-toxylem transfer. Such storage tissues buffer short-term and diurnal (17) fluctuations of nitrogen in both xylem and phloem.

In developing cottonwood shoots without secondary tissue, the metaxylem parenchyma cells are active in uptake and storage of both amino acids and sugars (Figs. 2-4). In the lower stem of cottonwood and in other plants with considerable secondary development, the ray parenchyma are the major uptake and storage tissues (5, 8). Sauter (25) has proposed that initial uptake from the vessels takes place in 'contact cells,' specialized ray cells active in loading and unloading the vessels. After uptake from the vessels, movement in rays results in both primary xylem-tosecondary xylem and xylem-to-phloem transfer. Autoradiographs of leaves and microautoradiographs of both leaf and stem transections from plants fed different amino acids or sucrose via the transpiration stream illustrate that each of the fed compounds has a somewhat different uptake (movement from the xylem free-space into the symplast) and distribution pattern. Such patterns are the product of the physical environment of the apoplast and the membrane transfer characteristics of the symplast.

Numerous studies have shown that xylem sap is slightly acidic (usual pH range of 5.0-6.0) and is maintained in this acidic state largely through the metabolism of malic acid and other strong organic acids (1, 9, 21). The pH of cottonwood xylem sap (5.3-5.6) is near the optimum found for alanine and glutamine uptake

Table VI. Distribution of ¹⁴C among Individual Organic Acids from Cottonwood Laminae and Stems after Xylem-Feeding Different Radioactive Amino Acids and Sucrose

The organic acids were separated from other water-alcohol-soluble compounds with ion exchange and then isolated with TLC. The different spots (corresponding to the location of standards run on the same TLC sheet) were scraped from the TLC sheets and counted with liquid scintillation spectrometry. Data shown are from the 1-h treatment. The origin contains sugar phosphates and other phosphorylated compounds that did not move with this solvent system. Threonine was not included because there was not enough ¹⁴C in the organic acids to count.

	Per Cent Total Organic Acids								
Organic Acids	Lam	ninae	Stems						
	LPI 4	LPI 8	LPI 4	LPI 9	LPI 12				
			%						
			Glutamine						
Origin	19	51	11	8	7				
Citrate	6	9	16	6	8				
Malate	15	10	32	54	68				
2-Ketoglutarate	6	4	10	12	4				
Succinate	49	20	24	18	12				
Fumarate	5	6	7	2	1				
			Glutamic aci	d					
Origin	11	12	5	5	5				
Citrate	5	5	6	4	6				
Malate	18	18	45	48	68				
2-Ketoglutarate	17	22	7	27	8				
Succinate	34	39	31	13	12				
Fumarate	15	4	6	3	1				
			Aspartic acid	1					
Origin	7	5	29	17	8				
Citrate	10	6	19	8	18				
Malate	63	74	26	39	36				
2-Ketoglutarate	7	3	8	7	18				
Succinate	12	10	16	24	18				
Fumarate	1	2	2	5	2				
			Alanine						
Origin	5	21	36	20	16				
Citrate	14	13	13	14	8				
Malate	49	29	22	36	19				
2-Ketoglutarate	6	14	5	4	27				
Succinate	11	14	22	24	26				
Fumarate	15	9	2	2	4				
			Sucrose						
Origin	40	49	23	23	24				
Citrate	14	18	22	22	28				
Malate	21	21	25	26	18				
2-Ketoglutarate	6	5	10	9	10				
Succinate	16	6	17	18	17				
Fumarate	3	1	3	2	3				
	•	-	-	-	-				

by tomato stems (30). The uptake of amino acids by cottonwood stems follows the sequence alanine > threonine > glutamic acid > aspartic acid (lower stem, Table I). This is the same general uptake pattern of basic > neutral > acidic amino acids found in tomato stems (30). The ionization state of the amino compound is probably not as important in uptake as membrane configuration, membrane affinity for a particular amino acid, and pH effect on trans-membrane carriers (11).

Sucrose uptake from the xylem free-space by the xylem parenchyma was so efficient in the lower stem, petioles, and major leaf veins that little of the applied sucrose reached the mesophyll of either mature or developing leaves. The rapid sucrose utilization by the vascular tissue of cottonwood and the restricted transfer to phloem was probably related to the sucrose concentration applied and contrasts with other xylem-feeding studies with sucrose. Sacalis and Durkin (22) applied [¹⁴C]sucrose in a 2% sucrose carrier to cut roses and found considerable ¹⁴Clabeled sucrose was transferred from xylem to phloem and moved up the stem to the flowers. In our work with cottonwood, no additional unlabeled sucrose was added to the artificial xylem sap. If the concentration of sucrose were increased (lower specific activity), more [¹⁴C]sucrose probably would be transferred from xylem to phloem in mature leaves for retransport in the phloem to developing leaves. This concentration effect may also be an important factor in the uptake and distribution patterns obtained with the amino acids. For example, alanine was heavily concentrated in the xylem parenchyma of the stem (Fig. 2C) with little movement into either mature or developing leaves. Van Bel *et al.* (31) found that low concentrations of alanine increased uptake rate in tomato stems. They suggested that alanine was concentrated and retained in parenchyma cells near the vessels at low input concentrations. At higher input concentrations, movement away from the supplying vessels would increase but uptake rate would decrease and more alanine would move through the system. Similar results were found for glutamine in cottonwood (Dickson, unpublished). All of the amino acids and sucrose in our studies were supplied at low concentrations and high specific activities, which would favor maximum uptake and retention by xylem parenchyma.

Three distribution patterns based on movement into leaves and uptake by stems were observed in cottonwood. (a) Alanine and sucrose were rapidly taken up and retained in the stem and major leaf traces with little movement to either phloem or developing leaves. Threonine and glutamine (5) were rapidly removed from the xylem free-space in the stem and transferred from primary xylem to secondary xylem, from xylem to phloem, then translocated in the phloem to developing leaves. (c) Aspartic and glutamic acid were less readily taken up in the stem and moved into the mature leaves. The movement of aspartic and glutamic acid into mature leaves provided a ready source of amino nitrogen for protein synthesis and maintenance in these leaves. In addition, phloem loading and retransport from these leaves would normally be directed to the lower stem and roots. Thus, aspartic and glutamic acid may be important in recycling amino nitrogen from shoots to roots in cottonwood.

Metabolic Conversion during Uptake and Distribution. Conversion to organic acids or incorporation into protein were the two major metabolic pathways followed by these xylem-applied amino acids (Tables III and IV). The rapid incorporation of threonine and alanine into protein is consistent with their accumulation in areas of developing tissue of both stems and leaves (Fig. 2). In addition, these amino acids were probably utilized in protein synthesis by mature parenchyma cells. Similar incorporation into protein of mature tissue was found in *Pisum* after xylem-feeding several different amino compounds (19). ¹⁴C from sucrose rapidly appeared in most of the major chemical fractions, indicating that its utilization was different from that of the amino acids. The large amount of ¹⁴C in the residue supports the microautoradiographic evidence for incorporation into developing cell walls (Fig. 4).

The distribution of ¹⁴C within the amino acid and organic acid fractions extracted from different plant tissues showed that conversion of one amino acid to another was limited and largely confined to compounds closely related to the fed compound (Table V). Glutamine was rapidly deaminated in both leaves and stems and the resulting glutamic acid converted to GAB. This was unexpected because the transport amides are resistant to metabolism (26). However, a similar conversion was reported for glutamine in pea (29).

Threonine metabolism was particularly interesting because it was incorporated primarily into protein with little apparent metabolism to other amino acids (Tables III–V). However, analysis of the individual amino acids of protein extracted from young developing cottonwood leaves (pronase extract, 6 N HCl hydrolysis, TLC separation) showed that more than 50% of the ¹⁴C from [¹⁴C]threonine was in isoleucine (Dickson, unpublished). Threonine contained most of the remainder of the ¹⁴C, although small amounts of ¹⁴C (up to 9%) were also found in valine. Threonine is readily converted to isoleucine (13). However, we did not analyze for isoleucine in the soluble amino acid fractions because it is not a major constituent of xylem sap (Table V).

All of the organic acids separated by TLC contained some ¹⁴C,

which indicates the rapid movement of ¹⁴C from these xylemfed compounds into the tricarboxylic acid cycle (13). Malic acid usually contained the most ¹⁴C but succinic acid, citric acid and 2-ketoglutaric acid also contained high levels. A similar distribution of ¹⁴C in organic acids was found in apple after feeding [¹⁴C]glutamic acid and [¹⁴C]aspartic acid to roots (27). Pyruvic acid, an important intermediate of alanine metabolism, was lost during extraction or TLC separation of organic acids. This may be part of the reason for the low content of ¹⁴C in the organic acid fraction after alanine treatment.

The rapid metabolism of these xylem-borne compounds in cottonwood was consistent with other xylem- (12, 26, 29) or root-feeding (27) studies. Although certain pathways predominate, the transport amino acids—glutamine, asparagine, glutamic acid, and aspartic acid—are major intermediates in nitrogen metabolism and provide direct access to the tricarboxylic acid cycle (13). Thus, the carbon and nitrogen supplied by these compounds can be transported throughout the plant and converted to almost any final product, depending on the current demands of the plant.

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